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to perform PCR on cells present in histochemical sections or cytochemical smears, e.g. for biological, forensic or pathological studies. The primer was one of a pair used to amplify papillomavirus DNA from human cervical cancer cells SiHa. A 449 bp PCR prod. was obtd. by this method whereas multiple primer pairs were needed for the same result using conventional PCR methods. See also AAQ34980-6. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cpds. N4-(1-4C alkyl) cytidine 5'-triphosphate (I) and N4-(1-4C alkyl)-2'-deoxycytidine 5'-triphosphate (II) are new. (I) and (II) serve as substrates for RA and DNA polymerases for incorporation of the N4-(1-4C alkyl)-cytosine moiety into oligo- and polymucleotides. They can be used in DNA sequence analysis, primer extension reactions and nucleic acid amplification. To assess the potential for using N4-methyl-dCTP in PCR amplification, reaction mixts. cong. the canonical nucleotide set were compared to mixts in which dCTP was replaced by the N4- methyl-tytosine analogue, in a PCR experiment designed to amplify a 293 bp sequence of HPV16 DNA. Using a high-temp. regimen the desired fragment was obtained regimen, conducted with AdTP or with N4-methyl-dCTP, A low-temp. regimen, conducted with dCTP or with N4-methyl-dCTP, A low-temp. mixt., cleanly produced identical amts. of the expected fragment as the sole amplification product. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New N4-alkyl-(deoxy) cytidine 5'-tri:phosphate cpds. - useful in DNA sequence analysis, primer extension reactions and nucleic acid amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N4-methyl-cytidine; N4-methyl-deoxycytidine; triphosphate; CTP; dc
substrate; polymerase; cytošine; oligonucleotide; polynucleotide;
sequence analysis; primer extension reaction; PCR;
polymerase chain reaction; amplifilication.
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                                                                                                                                                        Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HPV16/pT713 primer.
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29-SEP-1994
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Sequence 20 BP; 2 A; 2 C; 7 G; 9 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                An upstream primer (AAT05336) starting at base 1975 of the Arabidopsis thaliana peptide transport atptr2a gene (see AAT05334) and a downstream primer (AAT05337) starting at base 2528 were used in RT-PCR to measure the extent of ATPTR2A transcription in plant tissue. A 569 bp fragment the atptr2a open reading frame is generated
                                                                                                                                                                                                                                                                        Peptide transport gene, atptr2a, disease-resistance; fungus-resistance; insect-resistance; pathogen-resistance; herbicide-resistance; transgenic plant; crop improvement; polymerase chain reaction; primer; RT-PCR, ss.
                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Plant peptide transport genes - used to increase plant resistance to herbicidal peptide(s), pref. those produced by a plant pathogen.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; tive 0; Mismatches 2; Indels
 Length 20;
                              2; Indels
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Score 13.8; DB 1;
Pred. No. 8.6e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                              Peptide transport gene atptr2a PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Page 40; 79pp; English
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                                                           1308 CAAGACATACAACTACC 1324
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 Ouery Match
Best Local Similarity 88.2%;
Matches 15; Conservative (
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                                                                                                                                                                                                                   31-JAN-1996 (first entry)
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es 15; Conservative
                                                                                                                                                                                                                                                                                                                                               Arabidopsis thaliana.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-336935/43.
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pigment epithelium-derived factor (PEDF) has both neuronotrophic and gliastatic activity, making it useful in cases where neurons die quickly and glia tend to proliferate (gliosis), e.g. in CNS cell culture, in neurodegenerative diseases and in CNS injury. The neuronotrophic effect of PEDF is especially useful for enhancing survival of neuronal cell cultures intended for use in transplantation. These include cultures of human foetal brain cells and neural retina and photoreceptor cells. The gliastatic activity of PEDF can be applied to inhibiting glial cell cused for inhibiting PEDF can be used for inhibiting PEDF can be used for inhibiting PEDF can immunoassay for determining levels of PEDF in fluid, cellular or tissue samples e.g for determining againg and/or other degenerative diseases. Eight primers (AAT11661-68) were synthesised base on the cDNA sequence of PEDF and used to amplify fragments of the PEDF gene for later sequencing. Two primers (AAT11661, AAT11662) were used to amplify a 2 kilobase fragment from exon 3 to exon
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Pigment epithelium-derived factor, PEDF; neuronal cells; neurons; glial cells; gliastatic; gliosis; central nervous system; CNS; neurodegenerative disease; injury; neuronotrophic; brain cells; barkinson's disease; photoreceptor cells; retina; inhibition; proliferation; immunoassay; antibody; ageing; degenerative disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                          Use of pigment epithelium derived factor - for enhancing neuronal cell survival and inhibiting glial cell proliferation, useful, e.g. in CNS cell culture or to treat neuro-degenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Huntington's disease; animal model; transgenic animal; mouse; therapy; drug screening; mhd gene; polymerase chain reaction; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Pred. No. 8.6e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse Huntington's disease gene exon 5 primer P586.
                                                                                                                                                                                                                                                                                                                                                      Chader GJ, Becerra SP, Schwartz JP, Taniwaki T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                               (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 8; Page 38; 151pp; English
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94US-00367841.
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Best Local Similarity
                                                                                                                                                                                                                                                      07-JUN-1994;
                                                                                                                                               WO9533480-A1
                                                                                                                                                                                                                       06-JUN-1995;
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Neo-specific primer P8, (AAT78982), primer P586 (AAT78983) derived from exon 5 of the mouse Huntington's disease (HD) gene (see AAT78974), and primer P9 (AAT78984) derived from intron 5 of the gene were used in the genotype analysis of heterozygous transgenic mice embryos carrying a targeted mutation in exon 5. The results indicated that loss of function of the endogenous Hdh gene resulted in embryonic lethality during early post-implantation development. Transgenic mice can be used as models of HD
                                                                                                                                                                             Mouse Huntington's Disease gene - useful for generating transgenic mice as a model of Huntington's Disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fas; antibody; human; immunoglobulin; variable region; rheumatism; autoimmune disease; rheumatoid arthritis; therapy; CDR; heavy chain; complementarity determining region; PCR primer; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer SHR-16 for H chain of Fas specific antibody coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-Fas recombinant antibodies - useful for treating auto-immune diseases, especially rheumatoid arthritis.
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                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 20;
Pred. No. 8.6e+02;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                          Example 5; Page 31; 69pp; English.
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                                                                                              (UYBR-) UNIV BRITISH COLUMBIA
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Best Local Similarity
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                                        03-JUN-1996;
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              02-DEC-1996.
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Query Match RESULT 1049 Best Loca Matches AAT47350/ ID AAT4 à 셤

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protein
This sequence represents a primer for the coding sequence for the protein of the invention is a recombinant protein (A), that comprises at least one region corresponding to an immunoglobulin (Ig) variable region which enables the protein to recognise and specifically bind to an antigen, preferably human Fas, and has substantially no more immunogenicity in a human parient than a human antibody. The proteins are useful for treating autoimmune diseases, another the proteins are useful arthritis. (A) is based on a murine monoclonal antibody. As the protein lacks the constant region, it has substantially no more immunogenicity in the human patient than a human publicantially no more immunogenicity in the human patient than a human publicantially no more immunogenicity in the human patient than a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody
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Sequence 20 BP; 2 A; 8 C; 2 G; 8 T; 0 U; 0 Other;

Gaps ö 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; Live 0; Mismatches 2; Indels 1452 TCCATTCTTCCTCAGTC 1468 Conservative Local Similarity 15,

rccarrcrcrcrcrcrc 20

AAT47350 standard; DNA; 20 BP AAT47350;

(first entry) 10-SEP-1997

Variant #6 of universal primer sequence for M13mp18.

PCR; primer; amplify; polymerase chain reaction; bacteriophage; Ml3mpl8; cystic fibrosis transmembrane conductance regulator gene; multiplex PCR; chimeric primer; genetic screening; mutation detection; CFTR; Wilms Tumour gene; beta-thalassaemia gene; ss.

Synthetic.

WO9641012-A1

19-DEC-1996

96WO-US009637.

95US-00474450. 07-JUN-1995;

(GENZ) GENZYME CORP

Shuber AP;

WPI; 1997-052372/05.

Universal primer used for multiplex DNA amplification - allows simultaneous amplification of multiple DNA target sequences for high through-put genetic screening.

Claim 8; Page 10; 38pp; English.

AAT47345-T47374 represent variants of a universal primer sequence (see AAT47344) derived from the bacteriophage vector Ml3mpl8. This sequence can be used as half of the DNA primer of the invention. The primers are used for amplification of a target DNA sequence, and can be used in a multiplex PCR amplification. The primers have the sequence 5'-XY-3', where X is a sequence that does not hybridise to the target sequence (such as this sequence), and Y is a sequence contained within or flanking the target sequence. The melting temperature of a hybrid between X and its complement (in the absence of other sequences) is 60 degrees C. During early cycles of amplification, products are synthesised that contain the chimeric primers on either end. The primers then serve as high stringency recognition sequences for subsequent rounds of

ŏ amplification. As a result, the annealing efficiency of different primers and their targets in a multiplex amplification reaction is normalised, thereby reducing preferential amplification of certain targets. The chimeric primer comprises a 5' universal domain and a 3' target-specific domain. They are used for the simultaneous PCR amplification of multiple DNA targets in a sample. The primer containing AAT47344 is particularly useful in high-throughput genetic screening for detecting the presence of multiple defined targets e.g. to detect mutations in genes like the cystic fibrosis transmembrane conductance regulator (CFTR), the Wilms Tumour, and the beta-thalassaemia genes

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Sequence 20 BP; 4 A; 5 C; 8 G; 3 T; 0 U; 0 Other;

Gaps ö 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; iive 0; Mismatches 2; Indels Query Match
Best Local Similarity 88.2
Matches 15; Conservative

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RESULT 1050

AAV06254 standard; DNA; 20

22-APR-1998 (first entry) AAV06254;

Puromycin-sensitive aminopeptidase (PSA) antisense oligonucleotide 2.

psoriasis; Puromycin-sensitive aminopeptidase, PSA, treatment, cancer, psori proliferative disorder, hybridise, antisense oligonucleotide, ss

Homo sapiens. Synthetic.

WO9738114-A1

16-OCT-1997.

96WO-EP001518. 09-APR-1996;

96WO-EP001518 09-APR-1996;

(NOVS) NOVARTIS AG.

Tobler AR, Altmann K, Schlapbach Fontana A, Constam D,

WPI; 1997-512727/47

Isolated protein with puromycin-sensitive aminopeptidase activity - which may be used in treatment of proliferative disorders, including cancer and psoriasis.

Claim 36; Page 109; 141pp; English.

This antisense oligonuclectide is specifically hybridisable with selected DNA or RNA deriving from the puromycin-sensitive aminopeptidase (PSA)-99. This oligonuclectide is used for dignosing conditions associated with PSA expression. The human PSA-99 (875 amino acids) and the murine PSA-99 (970 amino acids) both exhibit PSA activity and can be used to generate anti-PSA artibodies. Cell lines which produce the antibody and host cells transfected with vector containing nucleic acid molecules encoding the PSA and the oligonuclectides can be used in assays for identification of agents which act by targeting PSA, for modulating PSA activity or agents which act by targeting PSA, for modulating PSA activity of endogenous PSA substrates, proliferation rate or viability of cells or induce apoptosis within cells by inhibiting PSA activity. Agents which induce apoptosis within cells by inhibiting PSA activity. Agents which can diminish PSA activity in cells we modulation of the amount of PSA in cells due to modulation of PSA synthesis, may be used in treatment of proliferative diseases, including tumours such as leukaemias and

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schultz621-3.rng

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0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; Live 0; Mismatches 2; Indels
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                                                                                           1308 CAAGACATACAACTACC 1324
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97US-0048740P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (WELL ) WELLCOME TRUST LTD. (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                        AAV85967 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                    10-FEB-1999 (first entry)
Query Match
Best Local Similarity 88.2
Matches 15; Conservative
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05-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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                                                                                                                                                                                                                                    AAV85967
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                                                                                                                                   셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences shown in AAV33257 to AAV33263 represent primers used for the PCR amplification of the human papillomavirus (HPV) type 16 genome contained in the human cervical cancer cell line SiHa. The invention provides a thermal cycler sample compartment optimised for holding and controlling he temperature of one or more microscopes which facilitates thermal cycling. It also contains a device (barrier) for protecting a microscope slide from fluid or vapour when the slide is sealed in the device, comprising a plastics material that has high thermal in the device, computivity, and is imperious to fluid or vapour, and is dimensioned so as to receive the slide. The new thermal cycling compartment is useful for performing in situ PCR for detection of target nucleic acid sequences directly from cells fixed onto a microscope slide, used in the field of cell biology, forensic science and clinical, veterinary and plant pathology. The modified heat blocks increase the speed and reliability of in situ PCR performed on microscope slides by accelerating and rendering more uniform the heat transfer which occurs during thermal cycling.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New thermal cycler for in-situ PCR on microscope slides - and device for protecting microscope slides from fluid or vapour.
                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human papillomavirus; HPV; human; cervical cancer cell line; SiHa; thermal cycler sample compartment; veterinary; thermal conductivity; in situ PCR; nucleic acid detection; PCR primer; ss.
                                                                                                                                        ;
0
                                                                                           Length 20;
                                                                                         Query Match
0.8%; Score 13.8; DB 1; Length 20
Best Local Similarity 88.2%; Pred. No. 8.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
carcinomas or epithelial disorders like psoriasis
                                            Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HPV type 16 gene amplifying 5' primer PV3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HOFF ) HOFFMANN LA ROCHE & CO AG F. (UNNY ) UNIV NEW YORK STATE RES FOUND.
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                                                                                                                                                                                     179 GAGGCATAGACAAGACC 195
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92EP-00306701.
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                                                                                                                                                                                                                                                                                                                                                    AAV33259 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
Human papillomavirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          correct PR field.)
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07-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP863213-A1
                                                                                                                                                                                                                                                                                                                                                                                                 AAV33259;
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TO AAV33259/C

AAV3259/C

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis; hypercholesterolemia; Alzheimer's disease; low density lipoprotein; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated LDL-receptor related protein - used to develop products for treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.
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Hey P, Kawaguchi Y, Merriman TR, Metzker ML, Nakagawa Y;
Phillips MS, Twells RCJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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rapidly cycle temperature of sample.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV69985 standard; DNA; 20 BP.
                                                                                                                                                                                               98WO-US004041
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Best Local Similarity 88.2%;
Matches 15; Conservative C
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                                                                                                                                                                                                                                                                                    Blumenfeld M, Chaplin J;
                                                                                                                                                                                                                                                         (MINU ) UNIV MINNESOTA
                                                                                                      Human papillomavirus.
                                                                                                                                                                                                                                                                                                                    WPI; 1998-495869/42.
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modified_base
                                                                                                                                                                                               03-MAR-1998;
                                                                                                                                                                                                                            03-MAR-1997;
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                                                                                                                                   WO9839479-A1
                                                                                                                                                                11-SEP-1998
                                                                                        Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The primers AAV43732-V43776 were to produce cancer associated gene fragments which can be used to detect cancer cells in tissue samples or biological fluids. They are detected by monitoring the change in mRNA expression as compared to normal tissue of one or more cancer-associated genes whose cDNA stringently hybridises to the nucleic acid fragments. The change in expression may be an increase or a decrease compared to normal tissue. The mRNA expression may be determined by PCR, Northern blotting or ribonuclease protection assay, or by determining the change in the amount of protein encoded by the gene(s) as compared to normal tissue, for example by using a labelled antibody recognising the protein. Detection of cancer cells for cancer diagnosis, including detection of metastatic cancer cells in tissues other than the primary tumour site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detection of cancer cells in tissue samples - by changes in mRNA expression compared to normal tissue of specific cancer-associated gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                       ss; cancer; PCR; Northern blotting; ribonuclease protection assay; diagnosis; metastatic cancer; primer; amplification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human papillomavirus (HPV) gene amplifying primer PV3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 2 A; 2 C; 7 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hino F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 67; 92pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Asada K,
                                                                                                                                                                                             Cancer associated gene primer 2.
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1435 GAGGATGCCATGAACA 1451
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                                                                                                      AAV43733 standard; DNA; 20 BP.
                 GAGGAGGCCATCAAACA 4
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                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mukai H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-467552/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yoshikawa Y,
                                                                                                                                                                                                                                                                                                                                                                                            21-FEB-1997;
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                                                                                                                                                               16-NOV-1998
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                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                  AAV43733;
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Sequences shown in AAV54677 to AAV54683 represent primers used for the PCR amplification of the Human papillomavirus (HPV) gene contained in the human cervical cancer cell line SiHa. These are used in the course of the invention which provides a thermal cycling device comprising a ceramic sample plate supporting a ceramic sample plate supporting a flat thermally coupled to the plate ample and a thermal sensor, a heater thermally coupled to the plate ample and a colorer for the substrate. The device atther maintains the temperature of the sample or subjects it to thermal cycling. The thin ceramic plate permits very rapid heat transfer to a sample on a microscope slide and this thermal cycling devise can be used for PCR amplification or hybridisation of target nucleic acid on microscope slide
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Human papillomavirus; HPV; thermal cycling device; ceramic sample plate; blological sample; thermal sensor; heater; cooler; thermal cycling; rapid heat transfer; microscope slide; PCR amplification; hybridisation; target nucleic acid; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thermal device for PCR amplification or hybridisation of target nucleic acid on microscope slide - has ceramic sample plate supporting flat substrate for sample and heater and cooler controlled to maintain or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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Pred. No. 8.6e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
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New nucleic acid encoding human cyclin-dependent kinase-10 - used e.g. to identify modulators of cell cycle progression for treating cancer or immune cell proliferation.
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                                                                                                                                                                                                                         Gene-specific primer PK22L234 and adapter primer API (see AAV32935) were used in a RACE PCR technique for cloning a 5' coding region of novel human cyclin-dependent Kinase 10 (CDX10) cDNA, using adapter-ligated human placenta cDNA as template. Nested primers (see AAV32936-37) were used in a second PCR, to produce an approximately 600 bp product. A 3' fragment was identified by database search, and a full-length sequence (see AAV32932) was produced in vector pLITMUS28:CDK10. The CDK10 protein product (see AAW3903) is used 6.g. to identify modulators of cell cycle progression and for treating cancer or immune cell proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New vector for plant transformation - useful for producing toxins that are specific to certain plants, or those which act on membrane systems and/or other cellular structures.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plant transformation, T-DNA, toxin; transgenic; transgenic food;
binary vector; PCR primer; barnase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20; 38.2%; Pred. No. 8.6e+02; v Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Barnase open reading frame fragment amplifying primer.
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Tigelaar H, Elzinga N;
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 5 A; 7 C; 5 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                             Example 1, Page 27; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1160 GGGGTGTGGCTGCATC 1176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
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Best Local Similarity 88.2%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGTCTGTGGGCTGCATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX05691 standard; DNA; 20
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Hoekstra S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-APR-1999 (first entry)
(MERI ) MERCK & CO INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-106063/09.
                                                                              WPI; 1998-447213/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stuiver MH,
Dekker BMM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9901563-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-JUN-1998;
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                                       Gerhold DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX05691;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV69978 to AAV69988 represent antisense oligonuclectides which are specifically hybridiaable with a region of a nucleic acid encoding human c-Jun protein. The antisense compound regulates the expression of the c-Jun protein. The present invention also describes antisense oligonuclectides which regulate the c-Fos protein. The antisense oligonuclectides which requisite the c-Fos protein. The antisense and isociated with Activating Protein lexpression, of which accompositions as c-Fos and/or c-Jun together with a carrier and a chemotherapeutic agent. They are used to regulate the expression of c-Fos and in calls or tissues, preferably by inhibiting metastasis. They also regulate cell cycle expression and can be used to regulate cell cycle expression and can be used to treat an animal with, or being prone to, a hyperproliferative disease
                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotides regulating Activating Protein 1 subunits - hybridise with c-fos and c-jun mRNA, used for regulating metastasis, cell cycle expression and hyperproliferative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CDK10; cyclin-dependent protein kinase; cell cycle; human; cancer; cell proliferation; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human cyclin-dependent protein kinase CDK10 cDNA primer PK22L234
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  /*tag= a
/note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 11 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                    Dean NM, Mckay R, Miraglia L, Baker B;
                                                                                                                                                                                                                                                                                                                                                                                                      Claim 12; Page 71; 120pp; English.
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97GB-00007491.
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                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                       WPI; 1998-609906/51
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                                                                                                                                   14-APR-1998;
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14-APR-1997;
                                                       WO9846272-A1
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cellular structures. Examples of genes include those encoding riboxymes against endogenous RNA transcripts, proteins evoking hypersensitive reactions, and RNA transcripts used for antisense/co- suppression inhibition of gene expression. The polymucleotide sequence contained the vectors prevents the transfer of DNA sequences beyond the T-DNA transgenic This avoids contamination of transgenic plants and/ or transgenic food with vector DNA. Sequences AAXO5690-91 represent primers used for the PCR amplification of the barnase open reading frame. This is used in the construction of a barnase expression cassette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention provides HIV cofactor inhibitors that contain oligonucleotides with a base sequence complementary to the CXCR4 or CCR5 genes. Such inhibitors can be formulated into drug compositions for prevention or treatment of HIV infection, with inhibition of expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of CXCR4 or/and CCR5 gene. Sequences AAZ31244-306 represent antisense oligonucleotides to the CCR5 gene
                                                                                                                                                                                                                                                                  Gaps
certain plants, or those which act on membrane systems and /or other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cofactor inhibitors, as drug
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inhibitor; HIV infection; CXCR4 gene; CCR5 gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 9.6e+02;
iive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                            Seguence 20 BP; 5 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CCR5 gene inhibiting antisense oligo AS(s)-60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Takai K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense oligonucleotide-based HIV cofactor compositions for treatment of HIV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yamamoto N, Kimura T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 6; Page 16; 59pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          drug composition; antisense; ss
                                                                                                                                                                                                                                                                                                    115 CCGATCGCCATGGATCG 131
                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ31303 standard; DNA; 20 BP
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Best Local Similarity 88.2.
Best Local Similarity 88.2.
                                                                                                                                                                                                                                                                  Conservative
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ31303;
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                                                                                                                                            Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genteal disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
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                                                                                                                 PCR primer used to amplify an ORF of Chlamydia trachomatis.
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Pred. No. 8.6e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genome sequence of Chlamydia trachomatis.
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97FR-00016034.
98US-0107077P.
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                            AAZ04231 standard; DNA; 20
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                                                                                      (first entry)
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Best Local Similarity 88.2
Matches 15, Conservative
                                                                                                                                                                                                                       Synthetic.
Chlamydia trachomatis.
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                                                                                                                                                                                                                                                                                                                                                                                                                     (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                               27-NOV-1998;
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04-NOV-1998;
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                                                         AAZ04231;
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AAZ02916
RESULT :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vaccine, eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; Inclusion conjunctivitis; genital disease; perhepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perhepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
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Pred. No. 8.6e+02;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                             Genome sequence of Chlamydia trachomatis.
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97FR-00016034.
98US-0107077P.
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Chlamydia trachomatis.
                                                                           Synthetic.
Chlamydia trachomatis
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Best Local Similarity
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17-DEC-1997;
04-NOV-1998;
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Matches
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PCR primers AA201426-206209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AA201425). These ORFs encode polypeptides (see AA20142494) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genteal diseases such as nongonococal urecritis, epidymitis, cervicitis, salpiniquis, perihepatitis, batholinitis; pneumopathy in breast feeding infants; and veneral lymphogranulomatosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The polypeptides of the invention may be of use in treating these
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                                                                                                                                                                                                                                                                                                                                                                                                         Genome sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 1754; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Deletion sequence oligonucleotide 2.
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                                                                                       97FR-00015041.
97FR-00016034.
98US-0107077P.
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                                                                                                                                                                                                                                                                                                                                            WPI; 1999-371125/31.
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                                                                                                                                                                                                                   (GEST ) GENSET
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                              27-NOV-1998;
                                                                                          28-NOV-1997;
                                                                                                                         17-DEC-1997;
04-NOV-1998;
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                                                                                                                                                                                                                                                                                Griffais R;
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This invention describes a novel composition comprising a number of sensor arrays, where each array comprises a unique probe oligonucleotide, which is the reverse complement of part of a unique target which is the reverse complement of part of a unique target on longonucleotides present in a mixture of target deletion sequence oligonucleotides. The compositions form a method for characterizing a sample of target deletion oligonucleotides which are labelled and hybridize with the probe oligonucleotides of the sensor arrays. Such oligonucleotides and their targets are represented in AAX2348-X3709. Oligonucleotides and their targets are represented in AAX2348-X3709. Oligonucleotides and their targets are represented in AAX2348-X3709. Oligonucleotides that are useful for modulating cellular addesion or proliferation, and being active against a enkaryotic pathogen, a human immunodeficiency virus (HIV), or a non-human retrovirus, including infilenza virus, Epstein Barr virus, Respiratory Syncytial Virus or cytomegalovirus (CWV). The compositions enable characterization of deletion sequence oligonucleotides having related, but different nucleobase sequences, and quantification of different specificity of the oligonucleotides in a mixture. Also, if the specificity of the oligonucleotides and may be performed using the secured of the method may be performed using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
oligonucleotides - useful for characterizing a sample of target deletion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'Match 0.8%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 8.6e+02; Hear 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 5 C; 5 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genome sequence of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 1480; Disclosure; 1912pp; English.
                                                        Example 1; Page 89; 163pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            133 ATGAAGAAGATCAAACG 149
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98US-0107078P
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                       oligonucleotides
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Matches
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ö AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX1990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent citis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX34584- AAX35879) can be used nucleotides sequences can also be used as immunogenic compositions as vaccines. Vectors containing C. pneumoniae epicope of C. pneumoniae The invention provides a DNA molecule comprising a sequence of mucleotides encoding an alphalF-subunit of a mammalian retinal calcium channel (RCC), including a human and antiha, a murine alphalF-subunit, a murine alphalF-subunit and orthologs of the human and murine alphalF-subunits. The RCC gene may be used to develop products for diagnostic tests, for incomplete CSMB and risk assessment in affected families. The RCC gene can provide information as to the basic defect in this retinal conditions, which could lead to effective methods for treatment or cure of the disorder. As the associated features of myopia, npstagmus and strabismus frequently observed in patients with incomplete CSNB may be caused by calciumtogluicated development pathways, identification of the RCC gene may help to elucidate the molecular details of eye development and which may lead to treatment for related eye disorders or diseases. Sequences AAZ46520-21 Retinal calcium channel, RCC gene, alphalF-subunit; retinal disorder, myopia, nystagmus; strabismus; calcium-regulated development pathway; eye disorder; human, EST; expressed sequence tag; CSNB; PCR primer; ss. New isolated mammalian retinal calcium channel gene, used to develop products for the diagnosis and treatment of incomplete congenital stationary night blindness and related disorders. Gaps ö 0.8%; Score 13.8; DB 1; Length 20; 88.2%; pred. No. 8.6e+02; ative 0; Mismatches 2; Indels Seguence 20 BP; 6 A; 6 C; 6 G; 2 T; 0 U; 0 Other; Human EST JRL4Al amplifying forward primer. Disclosure; Page 15; 55pp; English. (UYTE-) UNIV TECHNOLOGIES INT INC. 1468 CTGGGGGAGCGGATCCA 1484 20 99WO-CA000514. 98US-0087635P. CTGCGAGAGCGGATCCA AAZ46520 standard; DNA; 20 Bech-Hansen T, Naylor MJ; 13-MAR-2000 (first entry) Local Similarity 88.2 es 15; Conservative WPI; 2000-097327/08. 02-JUN-1999; 02-JUN-1998; Homo sapiens WO9963078-A2 09-DEC-1999. Synthetic AAZ46520; Query Match RESULT 1064 Matches AAZ46520 8866666666668888 ઠે 셤

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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at postition 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the pharmaceutical agents acting on a disease as well as other treatment. N. B. The SEQ ID NOS 2852, 2913, 3054, 3157, 3127, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel biallelic markers used to construct a high density disequilibrium
  represent primers for amplifying the human expressed seqeunce tsg (EST)
                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                               Human biallelic marker upstream amplification primer SEQ ID NO:4109.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human genome, biallelic marker, high density disequilibrium map, genomic map, haplotype, phenotype, polymorphic base, genotyping, haplotyping, in identification, characterisation, amplification, single nucleotide polymorphism, SNP, PCR primer, amplification, single nucleotide polymorphism, SNP, PCR primer,
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                                                                                          Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 8.6e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                        Sequence 20 BP; 1 A; 6 C; 2 G; 11 T; 0 U; 0 Other;
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                                                                                                                                                                    1698 TTACTCTGCCTACCT 1714
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                                                                                                                                                                                                                                                                                                AAZ69753 standard; DNA; 20 BP
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98US-0109732P.
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Best Local Similarity
Matches 15; Conserv
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23-NOV-1998;
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The invention relates to novel serine proteases designated BSSP6

(AABI1712-B11714), and to nucleic acids encoding them (AAA61763-A61765).

The invention also relates to vectors and transformants comprising BSSP6

nucleic acids; transgenic animals in which the expression level of BSSP6

can be varied; and an mBSSP6 knockout mouse. The invention additionally

encompasses anti-BSSP6 antichodies and methods of production of such

antibodies, methods of BSSP6 detection using the antibodies, and the use

of BSSP6 proteins or fragments as diagnostic markers for certain medical

conditions. Nucleotides encoding BSSP6 were initially isolated in a human

brain cDNA library using degenerate PCR primers (AAA6795-A61796) based

on conserved regions of serine proceases. The BSSP6 serine proteases and

nucleotides encoding them are useful in detecting homologues, mutants and

prostate gland, placenta, testis and spleen) as diagnostic markers for

conditions such as Alzheimer's disease, epilepsy, cancer, inflammation,

prostate pland, placenta, testis and spleen) as diagnostic markers for

conditions such as Alzheimer's disease, epilepsy, cancer, inflammation,

prostate pland prostatic hypertrophy. Sequences AAA61768-A61796

represent PCR primers used in the exemplifications of the invention.

Primers AAA61775-A61784 and AAA61795, while primers AAA61764-A61764

A61792 were used to isolate and annulify mutine BSSP6 cDNA (AAA61764).

Primers AAA61768-A61774 were used to construct plasmids used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Serine protease BSSP6, useful in detecting homologs, mutants and polymorphic variants as markers for diagnosis of Alzheimer's disease, epilepsy, cancer, inflammation, infertility and prostate hypertrophy, using blood or other tissues.
                                                                                                                                                                                                                                                                                                       BSSP6; serine protease; human; hBSSP6; mouse; mBSSP6; brain; diagnostic marker; antibody; transgenic animal; Alzhaimer's disease; epilepsy; cancer; inflammation; infertility; pancreatitis; prostatic hypertrophy; PCR primer; ss.
                                                                                                                                                                                                                                                                       Human serine protease BSSP6 (hBSSP6), RACE PCR primer, SEQ ID NO:23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yamaguchi N, Mitsui S;
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1060 ATCCCAACAAGACATA 1076
                                                                                                                                               ВЪ.
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                          18 ATCACAACACAGACATA
                                                                                                                                            AAA61782 standard; DNA; 20
                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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AAA61782/c
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Length 20; 2; Indels

Sequence 20 BP; 2 A; 7 C; 3 G; 8 T; 0 U; 0 Other;

Ouery Match 0.8%; Score 13.8; DB 1; Best Local Similarity 88.2%; Pred. No. 8.6e+02; Matches 15; Conservative 0; Mismatches 2;

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Gaps

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 /*tag= a
 /note= "Phosphorothioate linkages"

99US-00344001. 99US-00344001

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Location/Qualifiers

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The present invention describes an antisense compound (I) of 8-30 nucleobases, specifically hybridising to, and inhibiting expression of, human jun N-terminal kinase kinase-2 (JKK-2). Also described is a method of inhibiting the expression of human JKK-2 in human cells or tissues, comprising contacting the cells or tissues, with (I), in vitro. (I) has smitinflammatory, cytostatic and antinfectious activities, (I) is useful for inhibiting the expression of JKK-2 in human cells or tissues and prevents or delays infection, inflammation or tumour formation associated with altered expression of JKK-2 in human cells or detecting the levels of JKK-2 in a sample. The present sequence represents a phosphorothicate antisense oligonucleotide for human JKK-2, from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense compound specifically hybridizing and inhibiting the expression of human Jun N-terminal kinase kinase-2 is useful for treating infection,
                                                               Human, jun N-terminal kinase kinase-2; JKK-2; modulation; tumour;
antiinflammatory; cytostatic; antiinfectious; infection; inflammation;
detection; antisense therapy; phosphorothiaote; ss.
                                Human jun N-terminal kinase kinase-2 antisense oligonucleotide #33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Col 40; 31pp; English.
25-AUG-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammation and tumor.
                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-338506/29.
                                                                                                                                                                      Key
modified_base
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                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                per primers AAX89470-x89471 are used to screen a BAC library for the presence of a 14-3-3 sigma nucleotide sequence. 14-3-3 sigma nucleotide sequence. 14-3-3 sigma nucleotide sequence. 14-3-3 sigma nucleotide sequence. 14-3-3 sigma expression is regulated by p53 and exogenous expression of 14-3-3 sigma results in G2 block. The 14-3-3 sigma nucleotide and amino acid sequences are used in the invention to develop agents for the diagnosis, susceptibility determination and treatment of cancer. The amino acid sequence can be used in method for suppressing the growth of tumour centangence and be used for diagnosis, cells. The 14-3-3 sigma polypeptides can mediate cell cycle arrest upon damage to cellular DNA. 14-3-3 sigma probes can be used for diagnosing, testing cancers. They can also be used to treat other proliferative diseases, e.g. psoriasis, polyps, warts, and inflammatory diseases. The proliferation and crouse oligonucleotides can be used for promoting the
                                                                                                                                                                                                                                                    14-3-3 sigma; HWE1; stratifin; p53; diagnosis; cancer; psoriasis; polyp; psoriasis; wart; inflammatory disease; proliferation; ss; PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of 14-3-3 sigma polypeptides and nucleic acids for the diagnosis or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 20; 38.2%; Pred. No. 8.6e+02; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                     PCR primer used to screen a BAC library for 14-3-3 sigma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kinzler KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; Page 33; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proliferation and growth of cells
 9
                                                                                                                  AAX89471 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-0069416P.
98US-00210748.
                     20 GAGCACCAGAAGTGTGA 4
                                                                                                                                                                                                                                                                                                                                                                                                               98WO-US026924.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-022907/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
es 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hermeking H,
                                                                                                                                                                                                                                                                                                                                                                                                             18-DEC-1998;
                                                                                                                                                                                                                                                                                                                                            409931240-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                              18-DEC-1997;
                                                                                                                                                                                    15-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                           24-JUN-1999
                                                                                                                                                                                                                                                                                                           Synthetic.
 44
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Matches
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Gaps
                                                                                                                                                                                                                                                                                                                                                             Cytochrome P450; NADPH reductase; monooxygenase; CYP52A; CPR; POX; omega hydroxylase complex; omega-oxidation; fatty acid; alkane; alpha-omega-dicarboxylic acid production;
                                                                                     ;
0
                                                                                                                                                                                                                                                                                                                                  C. tropicalis CYP52A5A/CYP52A5B QC-RT-PCR primer, SEQ ID NO:47.
                                                     Score 13.8; DB 1; Length 20;
Pred. No. 8.6e+02;
0; Mismatches 2; Indels
                         Sequence 20 BP; 2 A; 4 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                    974 ACCGAGACCTCAAGCCC 990
                                                                                                                                                                                                                             BP.
                                                        0.8%;
                                                                                                                                                                                                                          AAA30532 standard; DNA; 20
                                                                                                                                                20 Acceceacercaaecec
                                                                                                                                                                                                                                                                                      (revised)
(first entry)
                                                                                       15; Conservative
                                                                       Local Similarity
present invention
                                                                                                                                                                                                                                                                                    15-SEP-2003
21-AUG-2000
                                                                                                                                                                                                                                                         AAA30532;
                                                                                                                                                                                               1069
                                                                                                                                                                                               RESULT 10
AAA30532
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Gaps

; 0

843 TGAGTACCTGGACAAGG 859

Conservative

18 reacracececacases

g

AAA29848 standard; DNA; 20

AAA29848

AAA29848/ ID AAA2 XX AC AAA2

schultz621-3.rng

AAA78243 standard; DNA; 20 BP.

RESULT 1070

AAA78243;

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The invention relates to 12 novel genomic DNA sequences and proteins which are components of the omega hydroxylase complex of Candida tropicalis ATCC 2036s. The DNA sequences (AAA30566-A3077) respectively encode cytochrome P450 NADPH oxidoreductases CPRA and CPRB (AAV30596, CYPS2ABA, CYPS2AB, CYPS2AB, CYPSZAB, CYPSZAB, CYPSZAB, CYPSZAB, CYPSZAB, CALABA, CALABA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fluids, agricultural chemicals, pharmaceuticals, dyes, surfactants, adhesives and fragrances. The CPR and CYR nucleic acids and proteins enable inexpensive large scale production of industrially useful dicarboxylic acids. Sequences AAA3D522-AD30543 represent QC-RT-PCR primers used in an exemplification of the invention to amplify the Candida tropicalis ATCC20366 CPR, CYP and beta-oxidation POX gene target mRNA.
quantitative competitive reverse transcription-PCR; QC-RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytochrome P450 nicotine adenine dinucleotide phosphate oxidoreductase and cytochrome P450 monooxygenase nucleic acids and encoded proteins, useful for overproducing dicarboxylic acids.
                                                                                                                                                                                                                                                                                                                                                                               Eirich LD, Eshoo M, Madduri KM, Cornett CA;
Loper JC, Gleeson M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ropicalis ATCC20366 CPR, CYP and beta-oxidation (Updated on 15-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 11; Page 44; 200pp; English.
                                              Candida tropicalis; ATCC20366
                                                                                                                                                                                                 99WO-US020797.
                                                                                                                                                                                                                                                  98US-0103099P.
99US-0123555P.
                                                                                                                                                                                                                                                                                                                                                                               Craft DL,
Tang M, 1
                                                                                                                                                                                                                                                                                                                              HENK ) HENKEL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-317711/27.
                                                                                                 WO200020566-A2.
                                                                                                                                                                                                   10-SEP-1999;
                                                                                                                                                                                                                                                  05-OCT-1998;
10-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                          Brenner AA,
                                                                                                                                                  13-APR-2000,
                                                                                                                                                                                                                                                                                                                                                                                 Wilson CR,
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Gaps ö ch 0.8%; Score 13.8; DB 1; Length 20; 1 Similarity 88.2%; Pred. No. 8.6e+02; 15; Conservative 0; Mismatches 2; Indels Query Match Best Local Similarity Matches 15; Conserv

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1010 AGAGGGGAGAGCTCAAG 1026
                                           2 AGAGGGCAGGGCTCAAG 18
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The present invention relates to antirheumatic agents which comprise as active ingredients an immunoglobulin M (IgM) protein. The IgM protein does not include a J segment, has apoptosis inducing activity, and consists of a light and heavy chain polypeptide produced synthetically. The agents of a light and heavy chain polypeptide produced synthetically. The agents of the invention exhibit antirheumatic and immunosuppressive activity and can be used to treat autoimmune diseases, especially rheumatism. The IgM molecule used in the invention has human Fas-antigen binding properties. Included in the invention are nucleotide sequences of the humanised anti-human Fas Ig CH11 (see AAA1222-A78212) and the corresponding protein sequences (see AAA12919-1812918 and AAA1202-A78212). And the corresponding protein sequences (see AAA1290-1812910). Also included are anti-human Fas antibody CD peptides (AAA1290-1812910). Primers specific for the anti-human Fas antibody, light, heavy and kappa chains used in the invention are represented by sequences AAA78313-A78318 and AAA78331-A78313, while humanised anti-human Ig DNA used in the invention are represented by sequences AAA78377-A78318 and AAA78331. Primer appresented by sequences AAA78377-A78318 and AAA78337. Primer represented by sequences AAA78377-A78318 and AAA78337. Primer represented by sequences AAA78371-A78333 and AAA78337. Primer represented by sequences AAA78337. Primer represented by sequences AAA78371-A78333 and AAA78337. Primer represented by sequences AAA78331-A78331 and AAA78337. Primer represented by sequences AAA783271-A78334 and AAA78337. Primer represented by sequences AAA78331. Primer represented by sequences AAA783271-A78334 and AAA78337. Primer represented by sequences AAA78331. Primer are used in the invention are represented by and are used in the invention are used in the inve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-human Fas humanizing antibody-containing antirheumatic agents.
                                                                                                                                          Anti-human Fas antibody CH11 H chain cDNA specific primer SHR-16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                        Antirheumatic agent; immunoglobulin M; IgM; apoptosis inducer; immunosuppression; autoimmune disease; treatment; rheumatism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; ve. 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 8 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the production of the agent of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 16; 109pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1452 TCCATTCTTCCTCAGTC 1468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EP.
                                                                                                                                                                                                                                             anti-Fas antibody; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   99JP-00263984.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98JP-00264598,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.2%;
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                                                                                             (first entry)
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Best Local Similarity 88.23
Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-454476/40.
                                                                                                                                                                                                                                                                                                                                             JP2000154149-A.
                                                                                             16-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                06-JUN-2000.
                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ59944;
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Lhx3; LIM-3; P-LIM; identification; characterisation; diagnosis; chromosome 9; pituitary disease; subtelomeric region; mutation; pituitary trophic hormone gene promoter; PCR primer; 88.

22-FEB-2000; 2000WO-US004424.

WO200050868-A2.

31-AUG-2000.

Homo sapiens

Human Lhx3 exon 6 PCR primer SEQ ID NO:113.

(first entry)

04-JAN-2001

AAA92148;

AAA92148 standard; DNA; 20 BP

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WPI; 2000-147195/13
                     22-JUN-1999;
                        22-JUN-1998;
21-JUN-1999;
            Homo sapiens
               W09967267-A1
                  29-DEC-1999.
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The introduction features Lie infallitication of genes whose expression tereins the cell with the drug of abuse, and comprise contacting the cell with the drug of abuse, and comprise contacting the cell with the drug of abuse, and detecting the expression of one or more of probes that specifically hybridise to the ESTs. The methods are used to identify of apuse, whose expression levels are altered by chronic or acute exposure to one or more drugs of abuse. The identification of genes whose regulation is altered in alcohol tolerance and/or addiction provides a valuable tool to evaluate the response to one or more drugs of abuse. Evaluation of the nature of this response provides information useful in designing to evaluate the response provides information useful in designing to evaluate the response provides information useful in designing to evaluate the response provides information useful in designing to evaluate and recovery regimens, and in evaluating the susceptibility of an organism or patient to drugs in a medical context. Monitoring the expression of identified genes and/or ESTs provides a mechanism by which tesponse of the organism to drugs of abuse. Sequences AAZSS941. Sequences AAZSS951 represent reverse transcriptase-PCR (RT-PCR) primers used to amplify a contal hybridisation probes from SH-SYSY-AH1861 cells in an expression in control and ethanol-treated SH-SYSY-AH1861 cells in an example dever dopamine beta-hydroxylase (DBH) and sodium-dependent corepinephrine transporter (NET). These genes whose expression and ethanol. Primers AAZS9944. and monocyte chemoattractant peptide 1 (MCP-1). These genes are thought to be important targets of ethanol. Primers AAZS9944. and monocyte chemoattractant peptide 1 (MCP-1). These genes are thought to be important beta-hydroxylase (DBH) probe The invention relates the identification of genes whose expression levels

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Gaps
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0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 8.6e+02;
tive 0; Mismatches 2; Indels
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400 GIGCAGICTCCAGIGAG 416

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18 GIGCAGIAGCCAGIGAG 2 셤

RESULT 1072 AAA92148

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Drug exposure; drug abuse; gene expression; EST; expressed sequence tag; identification; tolerance; addiction; therapy; screening; cellular response; ethanol; expression analysis; Northern blot; dopamine beta-hydroxylase; DBH; norephrephrine; reverse transcriptase-PCR; RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel methods for evaluating an organisms response to alcohol used to evaluate drug treatment and identifying susceptibility to alcohol.
Human dopamine beta-hydroxylase (DBH) PCR primer, SEQ ID NO:2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Page 68-69; 98pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Miles MF, Lai C, Lockhart DJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98US-0090268P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US013839
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Best Local Similarity 88.2°
Matches 15; Conservative
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The present invention describes an isolated nucleic acid (I) encoding a mammalian Lhx3. (I) is used in assays to: (I) detect and quantify the presence and level of expression of Lhx3. Lhx3a or Lhx3b, in a sample; (2) identify a compound that affects expression, the level of expression, or the activity of Lhx3, Lhx3a, or Lhx3b in a cell; (3) identify a normonant that affects binding or Lhx3 in a cell; (3) identify a bund of a pituitary trophic hormone gene promoter; (4) identify a human afflicted with a disease, disorder, or condition caused by altered and (5) detect a mutation in a Lhx3 allele in a human. The coding region of human Lhx3 has been genomically mapped to the subtelomeric region of human Lhx3 has been genomically mapped to the subtelomeric region of human Lhx3 has been genomically mapped to the subtelomeric region of human Lhx3 has been shown as P-LIM or LIM-3. The present sequence represents a PCR primer used in the amplification of human Lhx3, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated nucleic acid encoding mammalian Lhx3 for identifying a bwith a disease, disorder, or condition caused by an altered level of expression or binding of Lhx3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dog; genome; genomic marker; radiation hybrid map; identification; chromosome location; gene marker; polymorphic microsatellite marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dog genomic marker oligonucleotide sequence SEQ ID NO:746.
                                                                                                                                                                                                                                                                                                                                                                                                Rhodes SJ, Bridwell JL, Meier BC, Parker GB, Price JR;
Showalter AD, Sloop KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 2 A; 4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                  (ADRE-) ADVANCED RES & TECHNOLOGY INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 6; Page 169; 239pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA66884 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                               99US-0121110P
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Best Local Similarity 88.2%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                               22-FEB-1999;
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AAA66884
à
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clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5' and 3' ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the full length cDNA are useful length clones were obtained by construction of full length enriched cDNA. The full libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA enable the special methods. The present sequence is a primer used to amplify a human cDNA clone provided in the invention
                                                                                                                                     830 Primers useful for synthesizing full length cDNA clones and their use in genetic manipulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel transport system (A) for molecular substances (I) containing recombinantly prepared subunits (SU) based on amino acids (as) comprising: (i) at least two modified SU with one difference, and/or (ii) one or more modified SU with at least two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transport system for compounds, useful e.g. in gene therapy, comprises mosaic-like assembly of different protein subunits able to encapsulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Transport system; gene therapy; infection; tumor; ss; LLO; PCR primer; human immune deficiency virus; hemophilia; muscular dystrophy; capsid; cystic fibrosis; virus-like particle; cell targeting; listeriolysin O.
                                                                                                                                                                                                                                                        The invention relates to primers for synthesising full length cDNA
                        kawa T, Isogai T, Hayashi K, Ishii S, Kawai Y;
Sugiyama T, Nagai K, Kojima S, Otsuki T, Koga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 8.6e+02;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              L. monocytogenes listeriolysin O variant LLO PCR primer #3.
                                                                                                                                                                                                       Example 18; Page 132; 1380pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 7 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 11; Page 35; 106pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ď,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19 TGGACAGGAATGCAGAG 35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-NOV-2000; 2000WO-EP010876.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88.2%;
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AAH20451 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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nes 15; Conservative
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                          Nishikawa T,
                                                                                            WPI; 2001-524255/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-316433/33
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                          Ota T, Nishi
Wakamatsu A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH20451;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a radiation hybrid map of the dog (Canine familiaris) genome comprising the genome location of a marker selected from AAA66139 to AAA66942. The radiation hybrid map is useful for identifying and localising dog genes, since it covers approximately 80 % of the dog genome and provides a dense map integrating different types of the dog genome and provides a dense map integrating different types (i.e. Type I and Type II) of markers. The map and the dog genome markers for complementary sequences) are especially useful to identify genes responsible for phenotypic and behavioural traits in dogs, to identify morbid genes to analyse diseases and identify implicated genes in such diseases and their alleles, and to study dog pedigrees. They may also be unvolved in genetic diseases
                                                                                                                                                                                                                                                                                                                                                                                                          New radiation hybrid map of the dog, Canine familiaris, genome, useful for e.g. identifying genes implicated in phenotypic and behavioral traits or in genetic diseases and for studying dog pedigrees.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; full length cDNA; cDNA synthesis; oligo-capping; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cDNA clone-specific primer, SEQ ID NO: 4416.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
phenotype; behaviour; pedigree; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 85; 87pp; English.
                                                                                                                                                                                                                                                                           (CNRS ) CNRS CENT NAT RECH SCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             393 GGATGAGGTGCAGTCTC 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 GGAAGAGGTGCAATCTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAK95171 standard; DNA; 20 BP
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11-JAN-2000; 2000JP-00118774.
02-MAY-2000; 2000JP-00183765.
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Best Local Similarity 88.2
Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                       Andre C;
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-387821/33.
                                              Canis familiaris.
                                                                                          WO200029615-A2
                                                                                                                                                                                 15-NOV-1999;
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                                                                                                                                                                                                                             13-NOV-1998;
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RESULT 1074 AAK95171

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combined in a mosaic feashon to form (A) in which (I) can be encapsulated. (A) Are used to deliver (I) specifically to cells, particularly DNA to eukaryotic cells for gene therapy, e.g. of infections by human immune deficiency virus; tumors and a wide range of inherited diseases such as hemophilia, muscular dystrophy or cystic fibrosis. Capsids or other virus-like particles can be assembled, simply and in modular feabion, in vitro, allowing control over stoichiometric composition. SU can be modified to impart a wide variety of selected immunogenicity. (A) do not require extensive testing to ensure that they are safe (contrast replication-deficient viruses), also SU can be prepared in very pure form and are easily labeled fluorescently (for quality control or localization). This sequence represents a PCR primer used in the production of a Listeria monocytogenes listeriolysin LO variant which is used to illustrate the method of the invention
(optionally) unmodified SU. The various SU are
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 8.6e+02; Matches 15; Conservative 0; Mismatches 2; Indels
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differences; and (iii)
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AAH23201 standard; DNA; 20 BP. RESULT 1076

AAH23201;

17-SEP-2001

(first entry)

Human MMIF mRNA inhibiting antisense oligo ISIS #112711.

Macrophage migration inhibitory factor; MMIF; antisense; neurological; hyperproliferation; nootropic; antihormonal; immunosuppresive; human; antiinflammatory; cytostatic; ss.

Homo sapiens. Synthetic

WO200153317-A1

26-JUL-2001.

16-JAN-2001; 2001WO-US001475.

20-JAN-2000; 2000US-00489869.

(ISIS-) ISIS PHARM INC.

Murray SF, Cowsert LM, WPI; 2001-451899/48. New antisense compound(s) are useful to inhibit a nucleic acid molecule encoding macrophage migration inhibitory factor.

Claim 3; Page 82; 105pp; English.

The invention relates to antisense oligonucleotides 8-30 nucleotides in inhibitory factor (MMIF), where the antisense compound specifically hybridizes with and inhibitors he expression of MMIF. The antisense nucleotides with and inhibits the expression of MMIF. The antisense nucleotides are useful for the treatment of a disease or condition associated with MMIF such as neurological, hormonal, immune, inflammatory or hyperproliferative disorder. Sequences AAH2191-268 represent chimeric antisense phosphorothioate oligonucleotides used for inhibition of human MMIF mRNA expression

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The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acide acid. The immunostimulatory nucleic acides can be pyrimidine rich (py-tich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae) had over thomyvaviridae), bacterial antigens (e.g. toxophasma, and/or corthomyvaviridae), bacterial antigens (e.g. toxophasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Thi immune response and to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                                                                                                                      Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infections disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                       Gaps
                                                                     ;
0
                               0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 8.6e+02;
ative 0; Mismatches 2; Indels
Sequence 20 BP; 7 A; 4 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                      Immunostimulatory nucleic acid #929.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 101; Page 58; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vollmer J;
                                                               0;
                                                                                                           39 GGCAGGAGGACCAGCAG 55
                                                                                                                                             18
                                                                                                                                                                                                                                        AAF99813 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-SEP-1999; 99US-0156113P.
27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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(COLE-) COLEY PHARM GMBH.
                                                                                                                                             GGCAGAAGGACCAGGAG
                                                                                                                                                                                                                                                                                                              12-JUN-2001 (first entry)
                                                                       15; Conservative
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                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                            AAF99813;
                                                                                                                                                                                                       RESULT 1077
                                                                       Matches
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ö Gaps ö 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; ative 0; Mismatches 2; Indels Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other; 15; Conservative Best Local Similarity Matches δ

present sequence may have a phosphorothioate

1547 GCCTTCGGTCTTCGTCG 1563

GCCTTCGATCTTCGTTG 17

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ВР

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This invention describes novel regulatory sequences (A) derived from human fascin that provide specific expression in dendritic cells (DC) and which have antiviral, antibacterial, antibacatic, anti-
which have antiviral, antibacterial, antiparasitic, anti-
allergic, neurological, immunomodulatory and apoptocic activity. (A) are defensed to regulate expression of antigens, immunoregulators, antisense used to regulate expression of antigens, immunoregulators, antisense cells that contain (A) are useful: (1) in vaccines against viruses, bacteria, fungi, parasites, tumors, allergens and plaques in Creutzfeld-
actoria, fungi, parasites, tumors, allergens and plaques in Creutzfeld-
datoba and Alzheimer's diseases; and (ii) for gene therapy of tumors, allergies, infections, autoimmune diseases and transplant rejection. They can also be provide specific expression of antigens and immunoregulators in DC; for isolation and identification of capecific expression; to determine the degree of maturity of DC and to block transcription factoring binding sites in DC. (A) provide DC-specific expression; con contain and independent control, allowing a more specific expression of nucleic acid under their control, allowing a more specific expression of DC (since a complete leucoycte population may be transformed), including transformation in vitro. This sequence represents a primer associated with the human fascin gene described in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New regulatory sequences from the fascin gene, useful for providing dendritic cell-specific expression of e.g. antigens, e.g. for vaccination against tumors and infections.
                                                                                                                                                                                                         Fascin; regulatory sequence; human; dendritic cell; antiviral; tumor; antibacterial; antifungal; antiparastic; anti-allergic; neurological; immunomodulatory; apoptoclc; expression regulator; vaccine; allergen; Creutzfeld-Jakob disease; Alzheimer's disease; gene therapy; autoimmune disease; transplant rejection; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                  Human fascin associated primer SEQ ID 40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ross R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-JAN-2001; 2001WO-EP000362.
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02-MAR-2000; 2000DE-01010188.
                                    AAH48588 standard; DNA; 20
                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-451858/48
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(ROSS/) ROSS R.
(BROS/) BROS M.
                                                                                                                                                                                                                                                                                                                                                                                   WO200151631-A2
                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                           20-SEP-2001
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                                                                                AAH48588;
RESULT 1078
                      AAH48588/
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Bros M;

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The present invention relates to a retrovirus of type C morphology, which sediments in a sucrose gradient at a density of 1.16-1.18 g/l. The retrovirus is infectious for canine cells and belongs to the oncovirinae group. The present sequence is a PCR primer for the retrovirus of the present invention. The retrovirus can be included in pharmaceutical compositions or medicaments to treat autoimmune diseases, haematopoietic malignancies or malignant tumours, especially in humans. The retrovirus especially a human
                                                                                                                                                                                                     PCR primer; immunosuppressive; cytostatic; gene therapy; retrovirus; canine; autoimmune disease; haematopoietic malignancy; malignant tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell line, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human GABA(A) receptor-associated protein specific primer GABArp-R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New infectious retrovirus isolated from a canine cell line, useful producing medicaments to treat autoimmune diseases, hematopoietic malignancies or malignant tumors and in diagnosis and gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2; Indels
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8.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 13.8; I
88.2%; Pred. No. 8.6e
:ive 0; Mismatches
                                                                                                                                                                            Canine retroviral PCR primer MLVIN5700+.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 31; Fig 11; 131pp; English.
AGCCCCAGAACCTGCTC 1002
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              17 AGCCCCAGAACCCGCAC
                                                                                         AAC89128 standard; DNA; 20
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Best Local Similarity 88.27
                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                             Unidentified.
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986
                                                                                                                       AAC89128;
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                                                                               AAC891
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Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 8.6e+02; Matches 15; Conservative 0; Mismatches 2; Indels

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WO200119871-A2

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The invention provides an agent comprising a pyrone compound or dihydroxy compound of specified formulae given in the specification. The agent is used for correcting gene expression requlation errors. Errors in the following genes may be corrected: Li-6, Li-10, hemeoxygenase-1, prostaglandin G/H synthase-2, macrophage inflammatory protein-1-alpha, inflammatory protein-1-alpha, IL-7 receptor, macrophage inflammatory protein-1-beta, liver and activation-regulated chemokine, macrophage inflammatory protein-2-alpha, growth regulated chemokine, macrophage inflammatory protein-1, marrix metalloproteinase-9, migration inhibitory factor-related protein-1, marrix metalloproteinase-9, migration inhibitory factor-related protein-1, ranskerolase, adenosine AZa receptor, CD7 artigen properdin P factor. regulator of G-protein signaling-2, Nef-associated factor-1, myeloid cluckemia cell differentiation protein-1, signal peptidase complex, and also side-effects caused by them such as inflammation. Sequences AAH76220 contrast and a not the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ô
Pyrone; gene therapy; antiinflammatory; gene expression; interleukin; hemeoxygenase-1; prostaglandin G/H synthase-2; RANTES; TWF alpha; p78; macrophage inflammatory protein; chemokine; growth regulated protein-1; matrix metalloproteinase-9; migration inhibitory factor-related protein; lyzoxyme; GABA(A) receptor-associated protein; interferon; SCO homolog-2; lransketolase; adenosine Ala receptor; CD37 antigen properdin P factor; G-protein; Nef-associated factor-1; signal peptidase; PCR primer; se
                                                                                                                                                                                                                                                                                                                                                                                           Agent for correcting gene expression regulation error comprises pyrone compound or dihydroxy compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Heavy chain variable region; llama; Malassezia furfur; dandruff;
hair care; GAL7 promoter; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer used to amplify the left-hand GAL7 promoter region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                Kato I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 4 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                Nishimura K, Sagawa H,
                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 6; Page 77; 93pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                917 TGTTCCTGTTCCAGCTG 933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВР
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                                                                                                                                                                                                                                                    13-JAN-2000; 2000JP-00004989.
03-OCT-2000; 2000JP-00303711.
                                                                                                                                                                                                                      11-JAN-2001; 2001WO-JP000082
                                                                                                                                                                                                                                                                                                  (TAKI ) TAKARA SHUZO CO LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF80165 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15; Conservative
                                                                                                                                                                                                                                                                                                                                Enoki T, Yamashita S,
                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-514436/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                           WO200151480-A1.
                                                                                                                             Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-JUN-2001
                                                                                                                                                                                        19-JJ-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF80165;
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PCR primers AAF80165-66 were used to amplify the left-hand GAL7 promoter. The amplified sequence was used to express fusion proteins comprising a heavy chain variable region of an antibody isolated from llama, which was immunised with Malassezia furfur. M. furfur has been implicated in active agent, and used to produce a composition is conjugated to an active agent, and used to produce a composition for topical application, e.g. to the scalp. The topical composition, e.g. hair care products such as shampoos and conditioners, skin care lotions, shower gels, etc., is useful for targeting an active agent to a site at which M. furfur is present for the treatment and prevention of dandruff
                                                                                                                                                                                                                                                                                    Composition for use in targeting active agent, especially antimicrobial agent to scalp for treating, preventing dandruff, has active agent conjugated to antibody capable of binding specifically to Malassezia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphism; human; interleukin 4 receptor-alpha; IL4R-alpha;
allergic disease; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 8.6e+02;
artive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stephens JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 8 A; 3 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Duda A, Nandabalan K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human IL4Ralpha gene PCR primer #48.
                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; Page 33; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1472 GGGAGCGGATCCACAAA 1488
                                                                                                                                                                                                         Frenken LGJ, Van Der Vaart JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENA-) GENAISSANCE PHARM INC.
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                                        28-AUG-2000; 2000WO-EP008380.
                                                                                99EP-00307356
                                                                                                                                              (UNIL ) UNILEVER NV.
(HIND-) HINDUSTAN LEVER LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-APR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 88.2
les 15, Conservative
                                                                                                                      (UNIL ) UNILEVER PLC. (UNIL ) UNILEVER NV.
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                                                                                                                                                                                                                                                 WPI; 2001-257877/26.
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Windemuth AK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                16-SEP-1999;
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22-MAR-2001.
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WPI; 2001-103078/11
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The present invention relates to polymorphisms of the human interleukin 4 receptor-alpha gene (IL4R-alpha, see AAF57718 for the reference asquence). Polymorlecides comprising polymorphic gene variants are useful for therapeutic purposes. For example, where a patient may benefit from expression of a particular IL4Ralpha protein isoform, an expression desirable to decrease or block expression of a particular IL4Ralpha protein isoform, an expression of a patient. In may desirable to decrease or block expression of a particular IL4Ralpha isogene, which may be done by turning off by transforming a targeted organ, tissue or cell population with an expression vector that expresses high levels of untranslatable mRNA for the isogene. Specific therapeutics identified by these methods may be useful for allergic diseases. The present sequence is a PCR primer for human IL4R-alpha New isolated polynucleotide useful for the identification of therapeutics in allergic diseases is new. Sequence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other; Example 1; Page 61; 188pp; English.

Gaps ö ch 0.8%; Score 13.8; DB 1; Length 20; Similarity 88.2%; Pred. No. 8.6e+02; 15; .Conservative 0; Mismatches 2; Indels Local Similarity Query Match Best Loca Matches

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RESULT 1083

ABZ72182 standard; DNA; 20 BP.

ABZ72182;

03-APR-2003 (first entry)

Gene 216 SSCP detection primer SEQ ID NO 154.

Human, Gene 216; chromosome 20p13-p12; antiasthmatic; anorectic; antiinflammatory; gastrointestinal; gene therapy; vaccine; asthma; obesity; inflammatory bowel disease; primer; ss.

Synthetic.

WO200178894-A2.

25-OCT-2001

13-APR-2001; 2001WO-US012245

13-APR-2000; 2000US-00548797

(GENO-) GENOME THERAPEUTICS CORP.

Keith T;

WPI; 2001-639428/73.

Isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode, useful for the prevention, diagnosis and treatment of asthma, obesity and inflammatory bowel disease.

Example 10; Page 149; 520pp; English.

The invention relates to isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode. The mucleic acids and proteins may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate Gene 216 expression. For example, the nucleic acids (or vectors) and proteins may be used to treat disorders

cc associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of gene 216 by expressing canactive proteins or to supplement the patients own production of Gene 216 proteins. Additionally, the nucleic acids may be used to produce the secreted Gene 216 protein, by inserting the nucleic acids into a host complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids and complementary sequences may also be used as DNA probes in need of sequences in samples and therefore which patients may be in need of sequences in samples and therefore which patients may be in need of reforative therapy. The Gene 216 protein may also be used as antigens in the production of antibodies and activity. The anti-Gene 216 and in assays to identify and activity. The anti-Gene 216 antibodies may also be used as diagnostic and activity. The anti-Gene 216 antibodies may also be used as diagnostic gents for detecting the presence of Gene 216 proteins in samples (e.g. by enzyme linked immunosorbant assay or ELISA). Disorders that may be consented, diagnosed and/or treated by the above methods include, for example asthma, obesity and inflammatory bowel disease. The present conformation. The primers are used in the physical mapping of the gene (assay) polymorphism (SACP) analysis (ABZ72087-ABZ72184).

Conformational polymorphism and genotypling (ABZ72117-ABZ72184). \$

Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;

Gaps ö 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; ive 0; Mismatches 2; Indels 88.2%; 15; Conservative Query Match Best Local Similarity Matches 15; Conserv

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538 CCCATCTTTGACAAGCC 554 ო CCCTTCTGTGACAAGCC 13

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ABZ72122 Standard; DNA; 20 BP RESULT 1084

ABZ72122; ABZ72122

(first entry) 03-APR-2003

Gene 216 SSCP detection primer SEQ ID NO 94.

Human, Gene 216; chromosome 20p13-p12; antiasthmatic; anorectic; antiinflammatory; gastrointestinal; gene therapy; vaccine; asthma; obesity; inflammatory bowel disease; primer; ss.

Synthetic.

WO200178894-A2

25-OCT-2001.

13-APR-2001; 2001WO-US012245.

13-APR-2000; 2000US-00548797.

(GENO-) GENOME THERAPEUTICS CORP.

Keith T;

WPI; 2001-639428/73.

Isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode, useful for the prevention, diagnosis and treatment of asthma, obesity and inflammatory bowel disease.

Example 10; Page 149; 520pp; English

The invention relates to isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode. The nucleic acids and proteins

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may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate Gene 216 expression. For example, the nucleic acids for vectors) and proteins may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of gene 216 by expressing inactive proteins or to supplement the patients own production of Gene 216 proteins. Additionally, the nucleic acids may be used to produce the secreted Gene 216 protein, by inserting the nucleic acids into a host complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids and complementary sequences and therefore which patients may be in need of restorative therapy. The Gene 216 protein may also be used as antigens in the production of antibodies and activity. The anti-Gene 216 and in assays to identify and activity. The anti-Gene 216 antibodies may also be used as diagnostic agents for detecting the presence of Gene 216 proteins in samples (e.g. or the production of antibodies may also be used as diagnostic and activity. The anti-Gene 216 antibodies may also be used as diagnostic agents for detecting the presence of Gene 216 proteins in samples (e.g. by enzyme linked immunosorbant assay or ELISA). Disorders that may be prevented, diagnosed and/or treated by the above methods include, for example asthma, obesity and inflammatory bowel disease. The present sequence is that of a Gene 216 traded primer used in examples of the invention. The primers are used in the physical mapping of the gene invention and generolise (ABZ722081) analysis (ABZ722194), sequencing (ABZ72185-ABZ72268) and genetyping (Generalional polymorphism identification using single strand conformational polymorphism (Secol analysis (ABZ722194),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                      8499999999999999999999999
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Query Match

0.8%; Score 13.8; DB 1; Length 20;
Best Local Similarity 88.2%; Pred. No. 8.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels 538 CCCATCTTTGACAAGCC 554 CCCTTCTGTGACAAGCC 18 ò a

Gaps ö

> ABS71117 standard; DNA; 20 BP ABS71117; RESULT 1085 ABS71117/c

(first entry) 27-NOV-2002

Rat GPCR ligand Bv8 cDNA PCR primer RBv8-WR2.

G-protein coupled receptor; GPCR; ZAQ; human; ZAQC; ZAQT; rat; ZAQI; rZAQI; rZAQ2; mouse; I5E receptor; mI5E; GPR73; Bv8 protein; digestive disorder; central nervous system disorder; CNS; diarrhoea; bowel inflammation; constipation; food absorption disorder; nootropic; Alzheimer's disease; Parkinson's disease; schizophrenia; laxative; antiinflammatory; antidiarrhoeic; neuroleptic; neuroprotective; PCR;

Rattus sp.

WO200262944-A2.

15-AUG-2002.

01-FEB-2002; 2002WO-JP000852

02-FEB-2001; 2001JP-00026820,

(TAKE) TAKEDA CHEM IND LTD.

Watanabe T,

Masuda Y, Takatsu Y,

Ohtaki T, Hinuma S;

WPI; 2002-627537/67.

their ability to modify the binding of G-protein coupled receptor (GPCR) protein ZAQ and related proteins (human ZAQC, human ZAQT, rat ZAQI (TZAQI), TZAQZ, human and mouse ISE (MISE) receptor, and mouse GPR73) to their ligands (the mature form of human, mouse or rat BVB protein). The receptor protein and ligand are contacted in the presence or absence of the test compound. The compounds are useful in a drug composition for the treatment, and prevention of dispertive and central nervous system (CNS) disportion disorders, including bowel inflammation, diarrhoea, constipation, food absorption disorders, Alzheimer's disease, Parkinson's disease and schizophremia. The present sequence represents a PCR primer used in the Screening of compounds modifying the binding of G-protein coupled receptor protein ZAQ and related proteins to their ligands for use in present invention relates to a screening method for compounds for receptor protein ZAQ and related proteins to the treatment and diagnosis of digestive disorders. Example 5; Page 127; 197pp; Japanese. examples of the present invention

Sequence 20 BP; 3 A; 7 C; 3 G; 7 T; 0 U; 0 Other;

ö ö Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 8.6e+02; Matches 15; Conservative 0; Mismatches 2; Indels

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AAH77194 standard; DNA; 20 BP. (revised)
(first entry) 07-AUG-2003 24-JAN-2002 AAH77194; RESULT 1086

PCR primer PV3 used to amplify HPV in human cervical cancer cells.

Human; cervical cancer; human papilloma virus; PCR primer; PV3; SiHa; HPV; Thermal cycling; AIDS; 88.

Human papillomavirus.

US6300124-B1.

02-NOV-1999; 09-OCT-2001.

99US-00432012. 99US-00432012. 02-NOV-1999;

(MINU) UNIV MINNESOTA

Cibuzar GT, Schiller P, Arik Blumenfeld M, Bar-Cohen A,

WPI; 2002-009526/01.

Microscopic slide temperature control apparatus for medical diagnosis comprises coupling resistive heating element between the connection pads provided at opposing ends of slide.

Example 3; Col 27; 25pp; English.

The sequence represents PCR primer PV3. The primer was used in the invention to amplify DNA from cells of the stable human cervical cancer cell line SiHa, containing on integrated copy of human papilloma virus (HPV) type 16 per human genome. The invention relates to a novel thermal cycling device for regulating the temperature of a biological sample on filat substrate. The invention also includes an apparatus comprising the flat substrate for use in the thermal cycling device. The invention is

Best Loca Matches

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The invention relates to an isolated polypeptide, comprising a variant form of mouse or human SAC1 polypeptide. The variant form is associated with altered preference for carbohydrates, other sweeteners or ethanol. The polypeptide and its associated DNA sequence can be produced by recombinant techniques and is useful for preventing obesity, diabetes or alcoholism associated with SAC1 expression. The sequences are useful in careening for drugs and sweeteners. Recombinant cell lines and transgenic or repress function of SAC1. Predisposition to diabetes, obesity or alcoholism can be ascertained by testing any fluid or tissue of a human (such as blood, pancreas or tongue) for sequence variations of the SAC1 can be detected in a biological companient of the SAC1 locus may indicate a predisposition to diabetes, obesity and/or alcoholism and may provide a diagnostic mark. The polymucleotide can be detected in a biological cample by contacting the DNA with a probe to form a hybridisation complex which is then detected. The sequences represent cDNA encoding human and mouse SAC1 polypeptides and PCR primers specific for the SCA1 genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated polypeptide comprising variant form of mouse or human SACI polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism
                                                                                                                                                                                                         Human, mouse; SACI; carbohydrate; sweetener; ethanol; alcoholism; ss; obesity; diabetes; transgenic embryo; body tissue; body fluid; pancreas; blood; tongue; PCR primer; anorectic; antidiabetic; gene therapy;
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                                                                                                                                                               Murine SAC1 gene-specific oligonucleotide PCR primer #422.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chatterjee A,
Tordoff MG;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 14; Page 90; 239pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (WARN ) WARNER LAMBERT CO. (MONE-) MONELL CHEM SENSES CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            360 TGGGGAGAGTGACCAGG 376
                        BP.
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28-JUL-2000; 2000US-0221419P.
10-NOV-2000; 2000US-0247443P.
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                                                                                                                                                                                                                                                                                 protein replacement therapy.
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d DR, Ross D,
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                        AAS97855 standard; DNA; 20
                                                                                                                    (first entry)
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Ohmen JD, Reed DR,
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nes 15; Conserv
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                                                                                                                    12-MAR-2002
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                                                                     AAS97855;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides the protein and coding sequences of rice lesion formation inhibitor Sp17. The protein improves the heat stress of the plant, and can be used in the development of new breeds of plants for agriculture and horticulture. The present sequence is a PCR primer used to isolate the coding sequence of the invention
useful for medical diagnosis of diseases such as AIDS, also for amplification of nucleic acids in biological samples. The invention has the advantage that it enhances operatively as the heat resisting element is directly coupled to the microscopic slide, and reduces costs as the use of a heat sink is eliminated. (Updated on 07-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rice-originated gene, Sp17, that inhibits lesion formation and is applicable in improving heat stress of plants thus leading to prevention of lesion formation, for developing new breeds of plants for agriculture and horticulture.
                                                                                                                                                                                                                                                                                    Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rice; lesion formation inhibition; heat stress; agriculture; Sp17;
transgenic; plant; horticulture; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rice lesion inhibitor protein Sp17 coding sequence PCR primer #9.
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18.2%; Pred. No. 8.6e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                             ch 0.8%; Score 13.8; DB 1; Length 20; 1 Similarity 88.2%; Pred. No. 8.68+02; 15; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                     Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NAAG-) NAT INST AGROBIOLOGICAL SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 6; Page 47; 53pp; Japanese.
                                                                                                                                                                                                                                                                                                                              1308 CAAGACATACAACTACC 1324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             379 TCAGCCACGTCCTCGGA 395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                        19 CAAGACATACATCGACC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-OCT-2001; 2001WO-JP009153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAL46967 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yano M, Yamanouchi U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-372312/40
                                                                                                                                                                                                                                                           Local Similarity
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ABN89264;

RESULT 1088 AAS97855

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(first entry)

BP.

ABS78535 standard; DNA; 20

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The present invention describes an antisense compound (I), 16 to 30 bases in length targeted to specific base regions of a nucleic acid encoding human Talin. Also described. (a) an antisense compound up to 30 bases in length which inhibits the expression of human Talin; (b) a composition (II) comprising (I) or (a), and (c) inhibiting the expression of human Talin in human cells or tissues comprising contacting the cells or tissues in vitro with (I) or (a). (I) has antimicrobial, antiinflammatory and cytostatic activities, and can be used in antiesness gene therapy and as Talin expression inhibitor. (I) can be used: to prevent or delay infermantion or tumour formation, and in diagnostics, therapeutics prophylaxis, and in research reagents and kits. The present sequence represents a human Talin antisense chimeric phosphorothicate coligonic lection, inflammation or tumour formation.

Coligonic lection, and in research reagents and kits. The present sequence represents a human Talin antisense chimeric phosphorothicate at the 5' and 3' ends and a 10 nucleotide decoyy gap in the middle, which is used in an example from the present invention
                                                                                  Human, Talin, antimicrobial, antiinflammatory, cytostatic, inhibitor; antisense gene therapy, infection, inflammation, Talin inhibitor; tumour; antisense oligonucleotide; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound useful for inhibiting expression of Talin and for preventing or delaying infection, inflammation or tumor formation.
                                                   Human Talin antisense phosphorothioate oligonucleotide SEQ ID NO:77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-OCT-2000; 2000US-00702251.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-OCT-2000; 2000US-00702251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-470102/50.
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modified_base
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                                                                                                                                                               Homo sapiens
                 29-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-APR-2002
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Cowsert LM;

/*tag= a /mod_base= OTHER /note= "2'-methoxyethyl (2'-MOE) nucleotides" 16. .20

Location/Qualifiers

/*tag= c /mod_base= OTHER /note= "2'-methoxyethyl (2'-MOE) nucleotides"

The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first contenier housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour mestastasis, precancerous lession, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeods; Osler-Webber Syndrome, myocardial angiogenesis plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject. Anglogenesis inhibitor; ss; anglogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial anglogenesis; plaque meovascularisation; telangiectesia; haemophiliac joint; solgerosia; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar. Gaps ö Length 20; 2; Indels Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other; 0.8%; Score 13.8; DB 1; 88.2%; Pred. No. 8.6e+02; Angiogenesis inhibitory oligonucleotide #1019. 0; Mismatches Claim 2; Page 37; 276pp; English. GCCTTCGGTCTTCGTCG 1563 GCCTTCGATCTTCGTTG 17 .307/c ABK41307 standard; DNA; 20 BP. (COLE-) COLEY PHARM GROUP INC. 14-DEC-2001; 2001WO-US048458. 14-DEC-2000; 2000US-025534P. (first entry) 13-DEC-2002 (first entry) Conservative acid of the invention WPI; 2002-566690/60. Similarity WO200253141-A2. 21-MAY-2002 Bratzler RL; 15; 11-JUL-2002. Synthetic. ABK41307; ABS78535; 1547 Query Match Local RESULT 1091 Matches ABK41307, EXXXEX ઠે 셤

> ö Gaps .. 0 Score 13.8; DB 1; Length 20; Pred. No. 8.6e+02; 0; Mismatches 2; Indels 1571 ACTCAGGCAGGCCAGCT 1587 4 Acrereceaecearer 20 0.8%; Query Match
> Best Local Similarity 88.2.
> Perton 15; Conservative

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RESULT 1090

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The invention relates to a set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions on chromosomes 3, 10 and 19. The markers are useful for genotyping or estimating the frequency of an allele in a population, for detecting an association between a genotype or haplotype and a phenotype, e.g. a disoase involving drug responses, obesity or disorders related to obesity, such as hypernicaemia, digestive pathology, hepatic function disorders, cancer, cardiovascular disease, hypertension, hyperlipidaemia, insulin disorders, atheromatous disease and cardiac insulficiency. The markers are useful for detecting a statistical correlation between a biallelic marker allele and a phenotype and/or between a biallelic marker haplotype and a phenotype. Sequences ABK41305 and ABK41328-ABK41331 represent PCK primers used to amplify human LSR gene or USF2 gene biallelic markers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions on chromosomes 3, 10 and 19, useful, for e.g. detecting statistical correlations between marker allele and a phenotype.
                                        Human, obesity associated-biallelic marker; ss; LSR; USP2; PCR; primer; drug response, hyperuricaemia, digestive pathology; hypertension; cancer; hepatic function disorder; cardiovascular disease; hyperlipidaemia; insulin disorder; atheromatous disease; cardiac insufficiency; obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                         B,
                                                                                                                                                                                                                                                                                                                                                                                                                         Cohen D, Blumenfeld M, Chumakov I, Abderrahim H, Bihain
Human LSR gene biallelic marker upstream PCR primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 4 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 307; 311pp; English
                                                                                                                                                                                                                                                                                         28-JUN-2001; 2001WO-IB001477.
                                                                                                                                                                                                                                                                                                                                    18-JUL-2000; 2000US-0219704P.
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                                                                                                                                                                                                                                                                                                                                                                            (GEST ) GENSET.
                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                               24-JAN-2002.
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Gaps ; 0 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; rative 0; Mismatches 2; Indels 15; Conservative Query Match Best Local Similarity Best Loca Matches

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1417 CGAAATCGGATCTCCGC 1433 20 CGAAATAGGATCTCAGC ଚ 셤

ABQ93162 standard; DNA; 20 BP. (first entry) 29-AUG-2003 21-OCT-2002 ABQ93162; RESULT 1092 ABQ93162,

T. tauschii/wheat D genome microsatellite cfd60 right PCR primer.

Microsatellite marker; wheat; D genome; mapping; genotyping; polymorphism; phenotypic trait; OTL; quantitative trait locus; disease-associated gene; development factor; quality factor; resistence factor; wheat product; identification; detection; genetically modified wheat; PCR; primer; ss.

The invention relates to a map of the bread wheat D genome comprising the genome location of a microsatellite marker selected from a group of 185 cauch markers (ARQ9273-ARQ9217). The invention also encompasses the use of left (ARQ92918-ARQ93107) and right (ARQ93103-ARQ93237) primers to amplify and detect the microsatellite markers, and to identify genes a plenotypic trait of interest in wheat. Wheat is an allowatellotd species consisting of 3 diploid genomes designated A. B and of intendent of genome is thought to have been introduced in the constraint of the pecies. The D genome is thought to have been introduced in the most recent intercrossing between the amphipioid AABB and Triticum trauschiif (UD), probably involving only a limited number of genorypes of both species. Due to its polypioid genome, the large size of its genome, and its low level of polymorbing, the genetic mapping of wheat has to dark species. Due to its polypioid genome, the large size of its genome, and its low level of polymorbing, and are very polymorphic in length, between one and six nucleotides long, and are very polymorphic in length, propied sequences of perween one intercontent intercreatellites are codeminant, and exhibit Mandalian in dation, microsatellites are codeminant, and exhibit Mandalian in conservable into show little intraspectes polymorphic than the A or B genome. These microsatellite markers thus help to overcome some of the problems associated which is less polymorphic than the A or B genomes. These microsatellite markers thus help to overcome some of the problems associated with the genetic mapping of wheat. The wheat D genome problems associated primers and associated primers of the invention are useful for identifying genes and associated primers of the interest, most notably Qrus (quantitative trait loci). In particular they may be used for analysing genes and alleles implicated in factors conferring resistant or pathysing development factors; quality factors and factors conferring resistent and assess whether or Map of wheat D genome comprising the genome location of a microsatellite marker, useful for e.g. identifying genes responsible for a desired phenotypic trait, especially quantitative trait loci in wheat, and (INRG) INRA INST NAT RECH AGRONOMIQUE. Guyomarch H; Claim 4; Page 6; 105pp; English. 22-DEC-2000; 2000EP-00403659 22-DEC-2000; 2000EP-00403659 Bernard M, Sourdille P, WPI; 2002-550410/59. Aegilops tauschii. Triticum aestivum. EP1217079-A1 26-JUN-2002. diseases

0.8%; Score 13.8; DB 1; 88.2%; Pred. No. 8.6e+02; iive 0; Mismatches 2; Ouery Match Best Local Similarity 88.2' Matches 15; Conservative

Sequence 20 BP; 4 A; 3 C; 7 G; 6 T; 0 U; 0 Other;

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Gaps

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Length 20; 2; Indels

> CCTGCTCATCAACGAGA 1012 966

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Human; NOVX; NOV1; NOV2; NOV3; NOV4; NOV5; NOV6; NOV7; NOV9; NOV9), NOV10; NOV12; NOV13; NOV7.associated disorder; cardiomyopathy. atherosclerosis; diabetes; cancer; cell signal processing; AlDS; metabolic pathway; neuro-olfactory system disorder; neoplastic disorder; acquired immunodéficiency syndrome; inflammatory disorder; obesity; anorexia; cancer-associated cachexia; neurodegenerative disorder; immune disorder; graft versus host disease; Crohn's disease; multiple solteroals; haemophila; idiopathic thromboytopenic purpura; infectious disease; bacterial infection; fungal infection; RTO-PCR; ss; protozoal infection; viral infection; real time quantitative-PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Alsobrook JP, Tchernev V, Liu X, Spytek KA, Zerhusen B; Patturajan M, Grosse Wm, Lepley DM, Burgess CE, Shimkets R; Szekeres E, Vernet CAM, Li L, Casman SJ, Boldog F, Gorman L; Gangolli EA, Pernandes E, Rieger D, Edinger S, Gunther E, Millet I; Sciore P, Ellerman K, Macdougall J, Smithson G;
                                                                                                                                                                                                                                                                                                                                                                                                                                           21-DEC-2000; 2000US-0257495P.
22-DEC-2000; 2000US-0259171P.
20-FEB-2001; 2001US-0259191P.
22-MAR-2001; 2001US-0277826P.
22-MAR-2001; 2001US-0277826P.
11-APR-2001; 2001US-028981P.
13-APR-2001; 2001US-0283656P.
13-APR-2001; 2001US-0283656P.
10-AUG-2001; 2001US-0311754P.
                                                                                                                                                    Human NOV8 RTQ-PCR primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                    21-DEC-2001; 2001WO-US049519
20 CCTGCTCATCAAGTGA
                                                                 ABS51129 standard; DNA; 20
                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                             WO200250277-A2.
                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                         21-OCT-2002
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                                                                                              ABS51129;
                                         RESULT 1093
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New NOVX polypeptides and polynucleotides, useful in gene therapy, particularly for treating or preventing cardiomyopathy, atherosclerosis, diabetes, Crohn's disease, hemophilia or cancer in humans. WPI; 2002-508801/54.

Example 2; Page 254; 391pp; English.

The present invention relates to the isolation of novel human proteins referred to as NOVX, and the polynucleotide sequences encoding them. The NOVX proteins of the invention include NOVX-NOVY3 proteins, NOVX auscided and antibodies are useful for treating or preventing a NOVX-associated disorder, or alleviating a pathological state in a subject, atheroscierosis, diabetes, cancer (e.g. adenocarcinoma, lymphoma, prostate cancer, uterus cancer), disorders related to cell signal processing and metabolic pathways, disorders of the neuro-olfactory system (e.g. those induced by trauma, surgery and/or neoplastic disorders), acquired immunodeficiency syndrome (AIDS), inflammatory disorders (e.g. asthma) obesity, anorexia, cancer-associated cachexia, neurodegenerative disorders (e.g. Alzheimer's disease, Parkinson's

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an antisense oligonucleotide compound of 8 to 50 nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6 in cells or tissues. The oligonucleotides can be administered therapeutically or prophylactically to treat an animal having a disease or condition associated with caspase 6, such as Rieger's syndrome or ataxia telangiectasia, hyperproliferative disorder, a haematopoietic disorder, a bone metabolism or cholesterol disorder, various types of cancer, neurological conditions such as Alzheimer's disease and other deregulated apoptoric pathological conditions. This polynucleotide sequence represents a mouse caspase 6 oligonucleotide relating to the invention.

NOTE: This phosphorothioate oligonucleotide sequence has 2'-MOE wings and
disease), immune disorders, graft versus host disease, Crohn's disease, multiple sclerosis, haemophilia, idiopathic thrombocytopenic purpura, and infectious diseases (e.g. bacterial, fungal, protozoal or viral infections). The polymucleotide sequences are also useful in gene therapy. The present sequence represents a real time quantitative (RTQ)-PCR primer used in NOVX expression studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Muscular; cytostatic; nootropic; neuroprotective; ophthalmological; antilipaemic; osteopathic; caspase 6; Rieger's syndrome; bone metabolism; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder; haematopoietic disorder; cancer; neurological; Alzheimer's disease; apoptotic; mouse; murine; ds.
                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    An antisense oligonucleotide of 8 to 50 nucleotides in length that inhibits caspase 6, is useful for treating Rieger's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse caspase 6 antisense inhibition related oligo SEQ ID No 119.
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                                                                                                                                                                                        Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 8.6e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                  Sequence 20 BP; 10 A; 3 C; 5 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                               1240 TTCATCTTCCGTATCTT 1256
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                                                                                                                                                                                                                                       15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-471315/50.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL40400;
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schultz621-3.rng

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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in method comprises: (b) clones of the genomic libraries contained in multiwell plates independent of the marker as edgeded to the mixture to carry out an amplification reaction; (c) a signal oxresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to plates; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified products; (h) the clones in the multiwell plates are specified from the detected result, and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis, ABL42957 to ABL45322 represent per represent PCR primers for human chromosome and arrayed, The represent PCR primers for human chromosome 21q22.1, which are represent represent por human chromosome 21q22.1, which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; MEKK4 modulation; mitogen-activated protein kinase kinase 4; MTK1;
                                                                                                                                                                       Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human MEKK4 antisense oligonucleotide, ISIS #123107.
                                                                                                                        Human chromosome 1p36-35 PCR primer SEQ ID NO:752.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 19; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          479 CACTACCAGCTGACATC 495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 cacraccarcreacaec 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAD37172 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                  10-MAR-2000; 2000JP-00066716.
                                                                                                                                                                                                                                                                                                                                                                                   12-MAR-2001; 2001JP-00068285.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arraying genome clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-144136/19.
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                              PCR primer; ss.
                                                                                                                                                                                                                                                                                        JP2001321190-A.
                                                                                                                                                                                                                                              Homo sapiens.
                                                                           11-APR-2002
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                                                                                                                                                                                                                                                                                                                                        20-NOV-2001.
                              ABL43708;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a new antisense compound, comprising 8-30 nucleobases targeted to a nucleic acid molecule encoding human cytohesin-1, specifically hybridises with, and inhibits the expression of, human cytohesin-1, a guanine nucleotide exchange protein for ARP (ADP ribosylation factor). The antisense compound may be used in a pharmaceutical composition for inhibiting the expression of cytohesin-1 in human cells or tissues, and in treating a disease or condition associated with cytohesin-1 by administering to the human the antisense compound e.g. tumour or inflammation. The antisense compound e.g. tumour or inflammation. The antisense compound is also useful for diagnostics, therapeutics, prophylaxis and as research reagents and kite. The present sequence is an antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotide encoding human cytohesin-1, useful for preventing or treating a disease or condition associated with cytohesin-1 expression e.g. tumor or inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                               n, antisense, cytohesin-1, guanine nucleotide exchange protein; ARF; ribosylation factor; inflammation; antiinflammatory; tumour;
                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                       Human cytohesin-1 3' UTR antisense oligonucleotide, ISIS#111045.
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ch 0.8%; Score 13.8; DB 1; Length 20; Similarity 88.2%; Pred. No. 8.6e+02; 15; Conservative 0; Mismatches 2; Indel8
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88.2%; Pred. No. 8.6e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 1 A; 11 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bennett CF, Rothlein R, Kishimoto TK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
(BOEH ) BOEHRINGER INGELHEIM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; Page 81; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              733 GCACCCTGCACCGCCAT 749
                                                                                            211 CAGATAGGCCTGGATGA 227
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ABL43708
ID ABL43708 standard; DNA; 20 BP.
                                                                                                                                         3 cccacacaccrccarca 19
                                                                                                                                                                                                                                                             ABS73952 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-FEB-2001; 2001US-00791243.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       targeting human cytohesin-1
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                                                                                                                                                                                                                                                                                                                                                       (first entry)
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Best Local Similarity 88.2
Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200268584-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                         06-DEC-2002
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  Query Match
Best Local (
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                                               Matches
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Gaps

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MAP3K4; MAP three kinase 1; MAP/ERK kinase kinase 4; MAPKKK4; cytostatic; prophylaxis; immunological; hyperproliferative disorder; cancer; therapy; antisense; inflammatory; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound targeted to nucleic acid encoding mitogen-
activated protein kinase 4, useful for treating immunologic disorder,
inflammatory disorder or cancer.
                                                                                                                                                                                                                                                     note= "2'-methoxyethyl nucleotides"
                                                                                                                      note= "2'-methoxyethyl nucleotides'
                                                                                        'note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                   Ward DT, Gaarde WA, Monia BP, Wyatt JR;
                                                       Location/Qualifiers
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/*tag= c
/mod_base= OTHER
                                                                               base= OTHER
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16. .20
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                                                        Key
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The present invention relates to antisense compounds, compositions and methods for modulating the expression of MEKK4 (also referred as mitogencrivated protein Kinase kinase 4; MAPSK4; MAP FARE Kinase 1; MAP/ERK kinase kinase 4; MAPKKK4; MTK1). The antisense oligos are useful for inhibiting the expression of MEKK4 in cells or tissues. They are also useful for treating an animal having a disease or condition associated with MEKK4 such as immunological, inflammatory, hyperproliferative disorder or cancer. Sequences of the invention are also useful for They are also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human MEKK4 DNA. This sequence is used in the exemplification of the invention Claim 3, Page 92, 132pp, English

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20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
  Sequence
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Gaps

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843 TGAGTACCTGGACAAGG 859

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The present invention relates to a method of diagnosing a cellular proliferative disorder of breast tissue, which involves determining the state of methylation of one or more nucleic acids isolated from the subject, where the state of methylation of the nucleic acids as compared with a state of methylation of the nucleic acids as compared with a state of methylation of the nucleic acids as compared proliferative disorder of breast tissue is indicative of a cellular proliferative disorder of breast tissue in the subject. The nucleic acids may be TWHST, HOXAS, NBS-1, retinoic acid receptor beta (RAPBERA).

MIN-1 or RASSFIA. The method is useful for diagnosing and/or determining a predisposition to a cellular proliferative disorder, in particular proliferative disorder, in particular proliferative disorder, in particular proliferative disorder, in particular carcinoma, colloid carcinoma, tubular carcinoma, medullary carcinoma, metaplastic carcinoma, tubular carcinoma, medullary carcinoma, metaplastic carcinoma in situ. The present sequence is a primer used in the exemplification of the invention
                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Diagnosing and/or determining a predisposition to a cellular proliferative disorder of breast tissue, in particular breast cancer, by determining the state of methylation of one or more nucleic acids isolated from the subject.
                                                                                                                                                                                                                                                                                                                                                                                     Human; methylated gene; methylation; breast cancer; marker; WT-1; cell proliferative disorder; WWST; HOXA5; NES-1; RABbeta; cyclin D2; retinoic acid receptor beta; costrogen receptor; Wilms' tumour; 14.3.3 sigma; HIN-1; RASSFIA; tumour suppressor gene; hypermethylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Davidson N, Fackler MJ;
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88.2%; Pred. No. 8.6e+02;
tive 0; Mismatches 2; Indels
                                         2; Indels
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0.8%; Score 13.8; DB 1;
88.2%; Pred. No. 8.6e+02;
ive 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE.
                                                                                                                                                                                                                                                                                                                                                 Cyclin 14-3-3 sigma gene PCR primer #14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 46; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sukumar S, Evron E, Dooley WC,
                                                                                  150 GCAGCTGTCAATGACAC 166
                                                                                                                                                                                                                            ABT06434 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JAN-2002; 2002WO-US002455.
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                                                                                                                      18 GCAGTTGTCAAGGACAC
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Best Local Similarity 88.29
Matches 15; Conservative
                                         15; Conservative
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  Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying cells in which both alleles of a gene are modified, comprising modifying one allele by insertion or replacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are useful for identifying a gene that cells having both alleles of a gene are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that contributes to the virulence and/or pathogenicity of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a diploid fungus and for identifying a therapeutic agent for treatment of a mammalian activity of a gene product, preferably enzymatic activity, carbon cativity of a gene product, preferably enzymatic activity, carbon cativity. The method is useful for identifying a compound having the activity. The method is useful for identifying a compound having the ability to inhibit growth or proliferation of C. albicans cells and for primer used in the method of the invention. Note: The sequence is that of a proper or primer used in the method of the invention. Note: The sequence as the control of the invention is not represented in the present equence is that of a primer and a sequence is not represented in the primer because the primer because the control of the invention. Note: The sequence is that of a primer and a sequence is not represented in the present and primer because is that of a primer and a sequence is the present and a sequence and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or therapeutic intervention, by inactivating in the strain one allele of gene and placing other allele of the gene under conditional expression.
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                                                                                                                                                                                                                                                                                                                                                                 Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Constructing strains for identifying gene products as effective targets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 36; SEQ ID NO 5188; 167pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                         Candida albicans GRACE strain PCR primer SEQ ID NO 5188.
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                                                                                                                                                              ABZ30969 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-DEC-2000; 2000US-0259128P.
20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
26-DEC-2001; 2001WO-US049486.
                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ELIT-) ELITRA PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jiang B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-566694/60
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Candida albicans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200253728-A2.
                                                                                                                                                                                                                                                                   30-JAN-2003
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                                                                                                                                                                                                                ABZ30969;
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Ohlsen KL;

Bussey H,

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The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying cells in which both alleles of a gene are modified, comprising modifying one allele by insertion or replacement fragment with a heterologous promoter, so that expression of the second allele by the promoter, so that expression of the second allele is regulated by the promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells having both alleles modified are useful for identifying a gene that cells in which both alleles of a gene are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a diploid fungus and for identifying a therapeutic agent for treatment of a mammalian disease. (M1) is useful for identifying a compound which modulates the cativity of a gene product, preferably enzymatic activity, carbon compound catabolism, blosyntheric, transporter, transcriptional, transduction, DNA replication and cell division activity. The method is useful for identifying a compound having the ability to inhibit growth or prolifezation of C. albicans cells and for treating infection by C. albicans. The present sequence is that of a PCR primer used in the method of the invention. Note: The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                                                                                                                                  Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 36; SEQ ID NO 5598; 167pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                              Candida albicans GRACE strain PCR primer SEQ ID NO 5598.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bussey H,
1717 CTGAGCCATGTTCACCT 1733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boone C,
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                                                                                                                                                  ВР
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20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-DEC-2001; 2001WO-US049486
                                                                                                                                                ABZ31379 standard; DNA; 20
                           CTGAGCCTTGTGCACCT
                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ELIT-) ELITRA PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jiang B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-566694/60
                                                                                                                                                                                                                                                                                                                                                                                                              Candida albicans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200253728-A2.
                                                                                                                                                                                                                                       30-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-JUL-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Roemer T,
                                                                                                                                                                                           ABZ31379;
                                                                                                         RESULT 1100
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Gaps

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Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 8.6e+02; Matches 15; Conservative 0; Mismatches 2; Indels

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Gaps ., 0

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0; Mismatches

15; Conservative

Matches

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Novel isolated nucleic acid encoding cytochrome P450 and NADPH reductase enzymes of omega-hydroxylase complex of Candida tropicalis, useful for increasing production of dicarboxylic acids.
                                                                                                                                                                                                             CPRA; CPRB; CYP52AlA; CYP52A2A; CYP52A2B; CYP52A3A; CYP52A3B; CYP52A5A; CYP52A5B; CYP52A5B; CYP52A5B; CYP52A5B; CYP52A5B; CYP52A5B; CYP52A5B; CYP50CHCOME P450; NADPH2 reductase; omega-hydroxylase complex; dicarboxylic acid; 88; quantitative competitive reverse transcription PCR; QC-RT-PCR; primer.
                                                                                                                                                                                           Candida tropicalis CYP52A5A/CYP52A5B gene QC-RT-PCR primer 7581-97-F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Loper JC,
3 11:01:46 2004
                                                          1335 AGCCGAGGCCCTTTTGA 1351
                                                                       AGCCGATGCCCTTTGGA 17
                                                                                                                           ABK31851 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                      98US-0083798P.
98US-0103099P.
99US-0123555P.
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                                                                                                                                                                 (revised)
(first entry)
                                                                                                                                                                                                                                                             Candida tropicalis, 20336.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craft DL,
Tang M,
                                                                                                                                                                                                                                                                                                                                                                                   CRAFT D L.
EIRICH L D.
ESHOO M.
CORNETT C A.
CORNETT C A.
BRENNER A A.
TANG M.
LOPER J C.
GLEESON M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-138383/18.
                                                                                                                                                                                                                                                                                                                                                                              WILSON C R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brenner AA,
                                                                                                                                                                                                                                                                                US6331420-B1
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                                                                                                                                                                                                                                                                                                                                       01-MAY-1998;
05-0CT-1998;
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                                                                                                                                                                         23-APR-2002
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                                                                                                                                              ABK31851;
                                                                                                                                                                                                                                                                                                                                                                                               (EIRI/)
(ESHO/)
(MADD/)
(CORN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                            (TANG/)
(LOPE/)
(GLEE/)
Mon May
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                                                                                                         RESULT 1101
                                                                                                                  ABK3185:
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Madduri KM, Cornett CA; Eirich LD, Eshoo M, Loper JC, Gleeson M;

Example 11; Col 35-36; 173pp; English.

The present invention relates to the isolation of Candida tropicalis 20356 novel genes (CRRA, CPRE, CYP52A3A, CYP52A3A, CYP52A3B, CYP5A3B, CYP5ABA, CANDIDA, CANDIDA,

Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;

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Length 20;
 Score 13.8; DB 1;
Pred. No. 8.6e+02;
 0.8%;
Query Match
Best Local Similarity
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The invention relates to a nucleic acid encoding a adipose (ADP)
polypeptide which requiates, causes or contributes to obesity in an animal or a human. The polynucleotides, proteins, ant-adp antibodies, modulators of adp activity, adp antisense mucleic acids, expression vectors, adp transgenic animals are useful in the diagnosis and treatment of obesity, adiposites, bulimia, wasting (cachexia), eating disorders and/or disorders of body weight/body mass, weight loss due to cancer or infectious diseases, genetic disorders associated with hypogonadism e.g. Prader-Willi syndrome, laurence-Moon-Biedl syndrome, hypothyroidism, diabetes, Cushing's syndrome, endocrine disorders gastrointestinal diseases, inflammatory bowel disease, ulcerative colitis, and anorexia nervosa. They are also useful for treating disorders of body weight/mass creating inpans, and/or liposarcomas. The compositions are also useful for treating conditions associated with under weight e.g. enhancing or controlling fertility, controlling weight loss in acquired immunodeficiency syndrome (AIDS) or cancer patients. The present sequence is a PCR primer used to amplify an adp nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Adipose protein; se; adp; obesity; transgenic animal; obesity; adipositas; bulimia; wasting; cachexia; eating disorder; blosorder; b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               causes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel nucleic acid encoding adipose polypeptide which regulates, cause or contributes to obesity, useful for treating obesity, heart disease, hypertension, infertility, and controlling weight loss in cancer patients.
Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rothe M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dohrmann C, Haeder T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse adipose protein, adp, PCR primer #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 158; 188pp; English.
                                                      1010 AGAGGGGAGAGCTCAAG 1026
                                                                                                                   18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JUN-2000; 2000US-0211914P.
23-JUN-2000; 2000EP-00113049.
28-JUN-2000; 2000US-021451BP.
17-APR-2001; 2001EP-00109537.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUN-2001; 2001WO-EP006713
                                                                                                                                                                                                                                                                            ABK16359 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Broenner G, Ciossek T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (DEVE-) DEVELOGEN AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-106464/14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                      14-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-DEC-2001.
                                                                                                                                                                                                                                                                                                                                            ABK16359;
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AAD44838;

RESULT 1103

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The sequence represents a mouse syndecan-1 reverse transcription PCR primer. The invention relates to a novel use of a Smad3 inhibitor in preparing a medicament to treat or prevent wounds or fibrosis. The invention has antifibrotic and vulnerary activity. The Smad3 inhibitors are useful for preventing fibrosis and improving wound healing. The Smad3 antibodies, as reagents for research purposes, or the identification of other cellular gene products involved in the regulation of fibrosis and improvement of wound healing, as reagents in assays for screening for compounds that can be used in the prevention of fibrosis and improvement of wound healing, and as pharmaceutical reagents in protecting against fibrosis and improving wound healing related to Smad3. Compounds that that Smad3 may be used in inhibiting the activity of wild type and/or mutant Smad3 and in inhibiting the activity of wild type and/or smad3 and in identifying compounds that disrupt normal Smad3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of Smad3 inhibitors in preparing a medicament for treating or preventing wounds or fibrosis, or as reagents in assays for screening compounds for preventing fibrosis and improving of wound healing.
                                                                                                                                                                                                          Smad3; wound healing; fibrosis; antifibrotic; vulnerary; mouse; PCR primer; reverse transcription; syndecan-1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 8.6e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                 Mouse syndecan-1 reverse transcription PCR primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Roberts AB, Ashcroft GS, Russo A, Mitchell JB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example; Page 38; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1093 ACACTGTGGTACCGGCC 1109
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                                                                                                                                                                                                                                                                                                                                                                                                                        19-MAY-2000; 2000WO-US013725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAY-2000; 2000WO-US013725
                          ABA96039 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACACTGTGGAACCAGCC
                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                             WO200189556-A1.
                                                                                                                       08-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABQ66488;
                                                                                                                                                                                                                                                                                    Mus sp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to novel antisense oligonucleotides which are targetted to nucleic acids encoding human raf proteins and capable of inhibiting raf expression. The invention also relates to methods of inhibiting hyperproliferation of cells which involves contacting the hyperproliferating cells with a therapeutically effective amount of an oligonucleotide of the invention. The method is useful for treating cancer, anglogenesis or neovascularisation, especially ocular angiogenesis or neovascularisation. The present DNA sequence is human raf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Raf kinase; hyperproliferation; neovascularisation; ocular angiogenesis;
therapy; cancer; cytostatic; anti-angiogenic; vascular; ophthalmological;
antisense; ss.
                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating cancer, anglogenesis or neovascularization by administering antisense oligonucleotides targeted to human raf sequences.
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                                            0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Human raf kinase related antisense oligonucleotide #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 10 C; 0 G; 4 T; 0 U; 0 Other;
SQ . Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      kinase related antisense oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1152 TGACATGTGGGGTGTGG 1168
                                                                                                                                 867 GCAGTACCTGGATGACT 883
                                                                                                                                                                                                                                                                                              AAD44838 standard; DNA; 20 BP
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95US-00755806.
97US-00888982.
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98US-00143214.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94US-00250856.
                                                                                                                                                                 18 GGAGTGCCTGGATGACT
                                         Query Match
Best Local Similarity 88.23
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-597918/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1-MAY-1995
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Monia BP;

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Cytohesin-1; CT1; inhibit; cytostatic; antiinflammatory; cytostatic;
                                                            Human cytohesin-1 mRNA levels inhibitor #57.
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RESULT 1104

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Gaps

The invention relates to a novel antisense compound of 16-30 nucleotides targeted to any of 71 specified regions of the sequence that encodes human cytohesin-1 (CTI), where the compound hybridises and inhibits expression of human CTI. The compound of the invention has antiinflammatory, cytostatic, and anti-infective activity. The antisense compounds may have a use in antisense gene therapy. The antisense compounds are useful for treating or preventing disorders associated with expression of human CTI, e.g. infections, inflammation and tumours, and as research and diagnostic reagents Sequences ABQ66412-ABQ6651 represent chimeric phosphorothoate oligonucleotides, with 2'-WOE wings and a deoxy gap. The claimed sequences inhibit production of cytohesin-1 anti-infective; antisense gene therapy; infection; inflammation; tumour; human; ss; inhibitor. Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forenaic; environmental monitoring; food industry; feed industry; ss. New antisense compounds directed against human cytohesin-1, useful for treating and preventing infection, inflammation and tumors. Sequence 20 BP; 1 A; 11 C; 5 G; 3 T; 0 U; 0 Other; Capture oligonucleptide Zip ID#2505 oligo #9. Claim 14; Col 41; 40pp; English 733 GCACCCTGCACCGCCAT 749 04-APR-2001; 2001WO-US010958 14-APR-2000; 2000US-0197271P 30-OCT-2000; 2000US-00702246. 30-OCT-2000; 2000US-00702246 ABI95418 standard; DNA; 20 16-FEB-2002 (first entry) Local Similarity 88.7 Bennett CF, Cowsert LM; (ISIS-) ISIS PHARM INC. WPI; 2002-478385/51. 40200179548-A2 US6383809-B1 25-OCT-2001 07-MAY-2002 Synthetic. Synthetic ABI95418; Query Match RESULT 1106 Best Loca Matches ABI95418, ઠે g

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The present invention describes a method (M1) for designing capture oligonuclectide probes (1) for use on a support to which complementary oligonuclectide probes (11) will hybridise with little mismatch, where oligonuclectide probes (11) will hybridise with little mismatch, where (1) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Cryptococcus neoformans, Candida albicans and infectious agents e.g. Cryptococcus neoformans, Candida albicans and Salents in the same on parasitic infectious agents of special nearty virus and pollo virus, and parasitic infectious agents of selected from Onchoverva volvulus, Entamoba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 1 hydroxylase deficiency, Turner Syndrome and obesity defects.

CC medinesis. The method is also useful for detecting genetic diseases such as 1 hydroxylase deficiency, Turner Syndrome and obesity defects or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and fed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the persent oligonucleotide sequences. ABIS2074 to a presence or absence of the target nucleotide sequences. ABIS2074 to a the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, K-ras, PCR primer; probe, capture probe; mutation detection;
ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease;
infection; 21 hydroxylase deficiency; Turner Syndrome, obesity; cancer;
oncogene; tumour suppressor; human papillomavirus; forensit;
environmental monitoring; food industry; feed industry; ss.
                                                                                                                                              Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                              Kliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Capture oligonucleptide Zip ID#518 oligo #9.
                                                              Favis R,
                                                                                                                                                                                                                 Example 5; Fig 29; 300pp; English
                                                              Gerry NP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  288 ACTICGITCIGCACGG 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABI93431 standard; DNA; 20 BP
                  (CORR ) CORNELL RES FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18 AGTICGITCGGCACGGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-APR-2001; 2001WO-US010958.
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nes 15; Conservative
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                                                           Barany F, Zirvi M,
                                                                                                     WPI; 2002-034366/04
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0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; rative 0; Mismatches 2; Indels

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Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                                                  Example 5; Fig 29; 300pp; English.
    (CORR ) CORNELL RES FOUND INC.
                         WPI; 2002-034366/04
               Barany F,
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The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary coligonucleotide probes (II) will hybridise with little mismach, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents c.g. Salmon-lia, listeria monocytogenes and Haemophilus influenza, fungal confectious agents e.g. Crybtococcus neoformans, Candida albicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytorrophis cirus, Epsterin-Barr virus and pollo virus, and parasitic infectious agents calected from Onchoverva volvulus, Entamoba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 1 hydroxylase deficiency, Turner Syndrome and obesity defects.

CC method in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, cancer is specifically associated with a gene selected from BRCA1 gene, cancer is specifically associated with a gene selected from BRCA1 gene, cancer is specifically associated with a gene selected from BRCA1 gene, cancer is respected for environmental monitoring, forensics and the food and dentifying if ligation of the oligonucleotide probe sets occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABI32074 to CRA11974 by the present invention Length 20; Sequence 20 BP; 5 A; 5 C; 8 G; 2 T; 0 U; 0 Other; the present invention

Gaps .. 0 2; Indels 0.8%; Score 13.8; DB 1; llarity 88.2%; Pred. No. 8.6e+02; Conservative 0; Mismatches 2; Local Similarity tes 15; Conserv Query Match Best Loca Matches

567 cercercerereace 583 m 19 ccrccgrcgrgcaagcc g à

712/c ABL50712 standard; DNA; 20 (first entry) 19-JUN-2002 ABL50712; RESULT 1108

Rat G protein-coupled receptor protein PCR primer SEQ ID NO:67.

Rat; rZAQ1; rZAQ2; G protein-coupled receptor; GPCR; antidiarrheic; laxative; drug development; digestive organ disease; colitis; diarrhoea; constipation; malabsorption syndrome; diagnosis; gene therapy; PCR primer; ss.

Rattus sp.

WO200216607-A1.

28-FEB-2002

23-AUG-2001; 2001WO-JP007209.

24-AUG-2000; 2000JP-00253862.

(TAKE) TAKEDA CHEM IND LTD.

Shintani Y; Terao Y, WPI; 2002-269361/31.

Kliman R;

Favis R,

Gerry NP,

Zirvi M,

encoded DNAs, for developing drugs to treat diseases of the digestive organs, e.g. colitis, diarrhea, constipation and mal-absorption syndrome. and rat brain-originated G protein-coupled receptor proteins and

Example 5; Page 77; 135pp; Japanese.

The present invention describes human and rat brain-originated G protein-coupled receptor (GPCR) proteins. The GPCR sequences have antidiarrheic and laxarive activities. The GPCR sequences can be used for developing drugs to treat diseases of the digestive organs, e.g. colitis, diarrhoea, constipation and malabsorption syndrome, including gene diagnosis and therapy. The present sequence represents a PCR primer for rat GPCR, which is used in an example from the present invention

Sequence 20 BP; 3 A; 7 C; 3 G; 7 T; 0 U; 0 Other;

Gaps . 0 / Match 0.8%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Fred. No. 8.6e+02; tes 15; Conservative 0; Mismatches 2; Indels Query Match

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ABZ86270 standard; DNA; 20 BP. 17-OCT-2003 (first entry) ABZ86270; RESULT 1109 ABZ8627

Human oligonucleotide sequence.

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Human, antisense, lung dysfunction, nasal airway dysfunction, antiallergic, antiallamatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, cytostatic, gene therapy, antisense gene therapy, respiratory, lung, adenosine sensitivity, adenosine receptor; bronchodilation, bronchoconstriction, lung allergy, lung inflammation; respiratory disease; ds

Homo sapiens.

WO200285308-A2. 31-OCT-2002.

24-APR-2001; 2001US-0286137P. 23-APR-2002; 2002WO-US013135.

Katz E, Pabalan J, Aguilar D; S; Li Y, Sandrasagra A, Tang L, Shahabuddin (EPIG-) EPIGENESIS PHARM INC. Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating allments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Claim 15; SEQ ID NO 1512; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a

Page

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first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent composition of the invention can infillammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, but was obtained in electronic format directly from WIPO cat fip. wipo.int/pub/published_pot_sequences
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Seguence 20 BP; 4 A; 4 C; 5 G; 7 T; 0 U; 0 Other;

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                                                                                                                                             15; Conservative
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58 TGACTGCTGAAACCCAG 74 ۳ 19 rgacrgcrgaaaracag ద

Human oligonucleotide sequence. ABZ89410 standard; DNA; 20 BP. (first entry) 17-OCT-2003 ABZ89410; RESULT 1110 ABZ89410,

Human, antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy, respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Sandrasagra A, Katz E, L, Shahabuddin S; Li Y, San Tang L, Miller S, Nyce JW,

Aguilar D;

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 4652; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a

The invention relates to a novel pharmaceutical composition, which has a

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first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 6' or 3' end 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entinfilalmmatory steroid and ublquinone. A composition of the invention has antinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cust in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing levels of adenosine cofficent in a subject stissue, or treating bronchoconstriction, lung surfactant in a subject stissue, or treating bronchoconstriction, lung surfactant in a subject stissue, or treating bronchoconstriction, lote: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO cat fibs.
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Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 0.8%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 8.6e+02; es 15; Conservative 0; Mismatches 2; Indels
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first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antisathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cust in antisanse gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung sinflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fip. wipo.int/pub/published_pot_sequences
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0.8%; Score 13.8; DB 1; Length 20;
38.2%; Pred. No. 8.6e+02;
ive 0; Mismatches 2; Indels
                                                                    1444 ATGAAACATCCATTCTT 1460
                Local Similarity 88.2%;
les 15; Conservative
   Query Match
                    Best Loca
Matches
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3 ATGAAGCATCCATACTT

Human oligonucleotide sequence. ABZ91330 standard; DNA; 20 (first entry) 17-0CT-2003 ABZ91330; RESULT 1112 CXSXLTLLXAXXBXBXBXBXBXXBXXBXBXBXBXCCX

BP

Human, antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Katz E, Li Y, Sandrasagra A, K. Tang L, Shahabuddin S; Nyce JW, I Miller S,

Pabalan J, Aguilar

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

The invention relates to a novel pharmaceutical composition, which has a

Disclosure; SEQ ID NO 8608; 872pp; English.

Disclosure; SEQ ID NO 6572; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has

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first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antinflammatory steroid and ubiquinone. A composition of the invention has antinflammatory, antiallergic, antisthmatic, hypotensive, immunosuppressive, antiallergic, antisthmatic, hypotensive, immunosuppressive, antiallergic, antisthmatic, hypotensive, creventing a respiratory, lung or malignant disease or condition, also preventing a respiratory, lung or malignant creppiratory effect of an antial ammatory steroid in a subject, for reducing levels of adenosine receptor, producing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, broducing bronchodilation, increasing levels of adenosine receptor, broducing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung albergies, or a respiratory disease or condition. Note: The sequence data for this patent is not respresented in the printed specification, but was obtained in electronic format directly from WIPO at fig.
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Tang L, Shahabuddin S;
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Miller S,
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first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an initial ammatory steroid and ubiquinon. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition, is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antistral ammatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactent in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed of specification, but was obtained in electronic format directly from WIPO
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Sequence 20 BP; 2 A; 3 C; 9 G; 6 T; 0 U; 0 Other;

. . Match 0.8%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 8.6e+02; es 15; Conservative 0; Mismatches 2; Indels Query Match Best Loca Matches

548 ACAAGCCCCTCAGCCGC 564 18 ACAAGGCCCTCAACCGC 2 ઠે g

ABZ85750 standard; DNA; 20 ABZ85750; RESULT 1114

86

17-OCT-2003 (first entry)

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antializatinflammatory; antiallergic; antializathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchodilation; bronchodilation; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC. Katz E, Pabalan J, Aguilar D; S; Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Claim 15; SEQ ID NO 992; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a

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first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antisflammatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from Wipo at fipp. Wipo.int/pub/published_pot_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, retinal pigmented epithelium derived neurotrophic factor, PEDF; retinal disease, retinal tumour, retinoblastoma; retinal detachment, neuronal-retinal tumour; macular degeneration; retinitis pigmentosa; diabetic retinopathy; inherited and age-related pathology; tumour; ocular disease; nerve injury; serine protease related disorder; cytostatio; ophthalmological; antiinflammatory; antidiabetic; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating retinal disease such as retinal tumors, retinitis pigmentosa, macular degeneration and diabetic retinopathy, in a subject, involves administering Pigment Epithelium Derived Factor to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; Ltive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chader GJ, Becerra SP, Rodriguez IR,
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 6 A; 7 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human PDEF DNA, PCR primer 603.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      212 AGATAGGCCTGGATGAG 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-00894215.
92US-00952796.
94US-00279979.
95US-00377710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95US-00520373.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17 AGATGGGCCTGTATGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABS57272 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JAN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tombran-Tink J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US6451763-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-JUN-1992;
24-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUL-1994;
25-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Johnson LV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABS57272;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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The present invention relates to the isolation of a human retinal pigmented epithelium derived neurotrophic factor (FBDF), and polymucleotide sequences encoding it. The gene encoding human PEDF maps to chromosome 17p13.1-pter. The invention also describes a truncated version of PEDF referred to as PEDF-BH, vectors comprising nucleic acids encoding PEDF or PEDF-BH, and a method of using these sequences to treat retinal diseases such as retinal tumours (e.g. retinoblastoma), neuronal-retinal tumours, macular degeneration, retinitis pigmentosa, retinal detachment, diabetic retinopathy, inherited and age-related pathologies, tumours, ocular diseases, nerve injuries, and conditions resulting from the activity of serine proceases. The present sequence represents a PCR primer used to isolate human PEDF genomic clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method (M) for generating a substantially homogeneous population of undifferentiated cells (UC) from a biological sample (BS), which comprises subjecting BS or its sub-sample to tissue-disruption to provide a mixed population (MP) comprising UC, subjecting MP to a cell size-discrimination (SP) step, and simultaneously or sequentially with SD, subjecting the cell population obtained to a cell-surface marker-discrimination step. Also described: (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Generating substantially homogeneous population of undifferentiated cells from sample, by disrupting tissue sample, discriminating cells in population based on size and performing cell-surface marker-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuroprotective, antiparkinsonian; gene therapy, nervous system;
central nervous system; CNS; Alzheimer's disease; Parkinson's disease;
acute brain injury; CNS dysfunction; tissue regeneration; tissue repair;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               stem cell; NSC; undifferentiated; nootropic;
                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1;
38.2%; Pred. No. 8.6e+02;
ve 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse Emxl antisense PCR primer SEQ ID NO:66.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 10; Page 48; 90pp; English.
Example 48; Col 45; 53pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                          1631 CCAGCAGGCAGCGCTG 1647
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAAGCTGGCAGCGGCTG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-MAY-2002; 2002WO-AU000700.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUN-2001; 2001AU-00005403
                                                                                                                                                                                                                                                                                                                                                            1 Similarity 88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ80343 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Purification; neural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-140465/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200297067-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 discrimination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-MAY-2003
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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%X000000000000X8
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prepared by (M); (2) a composition (II) for use in cell replacement therapy, comprising a population of substantially homogeneous population of neural stem cells (NSCB) generated by (M); and (3) a composition (III) comprising a growth factor identified using a homogeneous population of NSCB generated by (M). (I) can have nootropic, neuroprotective and neighborkinsonian activities, and can be used in gene therapy. (M) is useful for generating a substantially homogeneous population of undifferentiated cells such as NSCB from a biological sample, and is useful for the replacement of neural tissue in a naimal.

(II) is useful in cell replacement therapy in an organ such as the brain or in the nervous system, preferably central nervous system (CNS), for treating a CNS disorder such as Alzheimer's disease, Parkinson's disease, centre brain injury and CNS dysfunction. (I) is useful for the repair or regeneration of tissue. ABZ80278 to ABZ80363 represent PCR primers which are used in an example from the present invention for markers defining
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human interleukin 12 p40 subunit antisense oligonucleotide ISIS #139149.
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                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; 88; antisense; interleukin 12 p40 subunit; antibacterial; antinflammatory; cytostatic; infection; inflammation; tumour.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "2'.methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                        Score 13.8; DB 1; Length 20;
Pred. No. 8.6e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                         Sequence 20 BP; 1 A; 10 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                        919 TICCIGITCCAGCIGCT 935
                                                                                                                                                                                                                                                                                                                                                                                                                        4 rrccrcrrccaccrrcr 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABX33976 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                        0.8%;
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Best Local Similarity 88.2'
....hes 15; Conservative
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-074100/07.
                                                                                                                                                                                                                                                            cell populations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified base
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Gaps

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Length 20; 2; Indels

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Best Local Similarity
                                                                                                                       US2003032607-A1.
                                                                                                                  Unidentified.
                                                                                            05-SEP-2003
                                                                                                                                      31-MAY-1994;
                                                                                                                                        31-MAY-1995;
26-NOV-1996;
       diagnostics
                                                                                                                            13-FEB-2003
                                                                                                                                             07-JUL-1997
                                                                                       ACD42154;
                                                        Query Match
                                                            Matches
                                                                                ACD42154,
                                                                             RESULT
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which is targeted to mRNA encoding human c-raf, a rraf or b-raf (raf is a protein kinase playing a regulatory role in signal transduction, required the mRNA encoding human c-raf, a rraf or b-raf (raf is a protein kinase playing a regulatory role in signal transduction, required the required required in language of included is a composition comprising the oligonucleotide and a pharmaceutically acceptable carrier. The antisense oligonucleotide is useful for inhibiting the expression of human raf in human cells or tissues, by contacting the human cells or tissues with the oligo. The oligo. is also is useful for treating or preventing a disease or condition associated with the expression of raf by administering it in combination with a chemotherapeutic agent to a human or cells of the human, where the expression of raf is abnormal expression, and the condition is a hyperproliferative condition such as cancer, angiogenesis or neovascularisation. The oligo. is also useful for inhibiting hyperproliferation of cells. The oligos are also useful as tools, for example for detecting and determining the role of raf expression in various cell functions and physiological processes and conditions and for diagnosing conditions associated with raf expression and for research purposes. The present sequence is an antisense oligonucleotide included in the sequence listing but not mentioned elsewhere in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Adenosine triphosphate (ATP)-binding cassette transporter subfamily C12;
                                                        invention relates to an oligonucleotide 8-50 nucleotides in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cystic fibrosis transmembrane conductance regulator; human; CFTR/MRP; multidrug resistance-like subgroup; somatic gene therapy; ABCC12; paroxysmal kinesigenic choreoathetosis; cysteinyl leukotriene; anionic drug; methotrexate; neutral drug; glutathione; glucuronate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 20;
18.2%; Pred. No. 8.6e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 6 A; 10 C; 0 G; 4 T; 0 U; 0 Other;
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(USSH ) US DEPT HEALTH & HUMAN SERVICES.
        Disclosure; Page 32; 42pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             anionic drug, methotrexate; n
sulphate conjugated drug; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 88.2
Les 15, Conservative
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                                                                                                                                                                          The invention relates to an antisense compound 20-50 nucleobases in length targeted to a start codon region, coding region, a stop codon region of a nucleic acid molecule encoding human interleukin 12 p40 subunit. The compound specifically hybridises with one of the regions and inhibits the expression of human Interleukin 12 p40 subunit. The new compound is useful for inhibiting the expression of human Interleukin 12 p40 subunit in cells or tissues and comprises contacting the cells or tissues in vitro with the compound, so that antisense compound may also be used as research reagents and diagnostics, and as treatment or prevention of disease states, e.g. to prevent or delay infection, inflammation or tumour formation, in animals and humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
New antisense chimeric oligonucleotide, useful for modulating the expression of human Interleukin 12 p40 subunit, in treating or preventing disease states in humans and animals, and as research reagents and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense; c-raf; a-raf; b-raf; protein kinase; cancer; ss; signal transduction; cell proliferation; lung carcinoma; cytostatic; antisense gene therapy; chemotherapeutic agent; angiogenesis; hyperproliferative condition; neovascularisation; ocular angiogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is an antisense oligonucleotide of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human raf-associated antisense oligonucleotide #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                               Example 15; Col 45; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            480 ACTACCAGCTGACATCC 496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 Actoccadoridactic 19
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96US-00756806.
97US-00888982.
98WO-US013961.
98US-00143214.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-JAN-2002; 2002US-00057550.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94US-00250856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-FEB-2000; 2000US-00506073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACD42154 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
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New ATP-binding cassette transporter gene subfamily C12, ABCC12 polypeptide, useful for preventing or treating paroxysmal kinesigenic

Dean M;

Denefle P,

Rosier-Montus M, Prades C, Arnould-Reguigne I, Allikmets $R_{\it i}$

WPI; 2003-093101/08.

Novel antisense oligonucleotide which is targeted to mRNA encoding human raf and which is capable of inhibiting raf expression, useful for treating or preventing hyperproliferative conditions such as cancer.

WPI; 2003-503332/47.

Monia BP;

06-JUL-1998;

28-AUG-1998

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Gaps

chorecathetosis

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This invention describes a novel human ABCC12 (adenosine triphosphate (ATP) binding cassette transporter gene subfamily C12, i.e. cystic fibrosis transmembrane conductance regulator/multidung resistance-like subgroup (CFTR/MRP) family) polypeptide and its encoding polymucleotides ABCC12 polypeptide is useful for screening agonists and antagonist of the ABCC12 polypeptide. The products of the invention are useful for screening and active ingredient for preventing and treating paroxysmal screening an active ingredient for preventing and treating paroxysmal kinesport of organic anion transporters such as cysteinyl leukotriene, anionic drugs, such as methotrexate, neutral drugs conjugated to acidic ligands, such as glucathione, glucuromate or suphate conjugated drugs and can be used for somatic gene therapy. This sequence represents a region corresponding to an exon/intron boundary from the gene encoding a human ABCC12 isoform described in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, mouse, 88; primer, gene 216; antiasthmatic, antiinflammatory, anorectic; chromosome 20p13-p12; single nucleotide polymorphism; SNP; gene therapy, respiratory disease; asthma; obesity; PCR; bronchial hyper-responsiveness; chronic obstructive pulmonary disease; adult respiratory distress syndrome; inflammatory bowel syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated gene 216 nucleic acids, useful for diagnosing, preventing treating a disorder, such as asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease, obesity or inflammatory bowel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human gene 216 polymorphism detection PCR primer #32.
                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Little RD, Van Berdewegh P,
Allen K, Pandit S;
                                  Disclosure; Page 44; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SCHERING CORP.
GENOME THERAPEUTICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                 865 AAGCAGTACCTGGATGA 881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-APR-2002; 2002WO-US012063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-APR-2001; 2001US-00834597.
13-APR-2001; 2001WO-US012245.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19 AGGCATTACCTGGATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABX74975 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-092960/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200283077-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Keith T,
Simon J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABX74975;
                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SCHE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1120
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(first entry)

BP

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identified from human chromosome 20p13-p12. The invention also discloses regions of the 216 gene that contain single nucleotide polymorphisms (SNP's) which may be used as markers for disease susceptibility or severity. The nucleotides of the invention may have antiasthmatic, antinflammatory or anoxettic activities and may be used in gene therapy. The nucleotides of its fragments are useful for diagnosing, preventing or treating a disorder, such as respiratory diseases (e.g. asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease or adult respiratory distress syndrome), obesity, or inflammatory bowel syndrome. The nucleic acids are also useful for identifying increased susceptibility of a subject to the disorders mentioned. The nucleic acids are also useful for identifying conclear acids can also be used as primers and templates for the recombinant production of disorder-associated peptides or polypeptides, for chromosome and gene mapping, or for tissue distribution studies. The present sequence represents a gene 216 specific PCR primer used in the ö New isolated gene 216 nucleic acids, useful for diagnosing, preventing or treating a disorder, such as asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease, obesity or inflammatory bowel This invention relates to a novel isolated nucleic acid, gene 216, identified from human chromosome 20p13-p12. The invention also discloses regions of the 216 gene that contain single nucleotide polymorphisms Human, mouse, ss; primer; gene 216; antiasthmatic; antiinflammatory; anorectic; chromosome 20p13-p12; single nucleotide polymorphism; SNP; gene therapy; respiratory disease; asthma; obesity; PCR; bronchial hyper-responsiveness; chronic obstructive pulmonary disease; adult respiratory distress syndrome; inflammatory bowel syndrome. Gaps Del Mastro RG; ô DB 1; Length 20; 2; Indels Human gene 216 polymorphism detection PCR primer #92. Little RD, Van Eerdewegh P, Dupuis J, Allen K, Pandit S; Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other; Score 13.8; DB 1; Pred. No. 8.6e+02; 0; Mismatches 2 Example 10; Page 155; 650pp; English. (SCHE) SCHERING CORP. (GENO-) GENOME THERAPEUTICS CORP ö 538 CCCATCTTTGACAAGCC 554 cccrrcrcrdachadcc 18 ABX75035 standard; DNA; 20 BP. 0.8%; 15-APR-2002; 2002WO-US012063. 13-APR-2001; 2001US-00834597, 13-APR-2001; 2001WO-US012245. 25-MAR-2003 (first entry) Local Similarity 88.2 18s 15; Conservative scope of the invention WPI; 2003-092960/08 WO200283077-A2. Homo sapiens. 24-OCT-2002. Keith T, Simon J, syndrome. Query Match ABX75035; RESULT 1121 Matches 888888888888888888888888888888888888 ò 셤

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Del Mastro RG;

Dupuis J,

This invention relates to a novel isolated nucleic acid, gene 216,

Example 10; Page 155; 650pp; English.

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Gaps

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Length 20; 2; Indels

0.8%; Score 13.8; DB 1; 88.2%; Pred. No. 8.6e+02; 1ve 0; Mismatches 2;

88.2%;

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(SNP's) which may be used as markers for disease susceptibility or antinflammatory. The nucleotides of the invention may have antiasthmatic, antinflammatory or anorectic activities and may be used in gene therapy. The nucleic acids, antibodies or its fragments are useful for diagnosing, preventing or treating a disorder, such as respiratory diseases (e.g. preventing or treating a disorder, such as respiratory diseases (e.g. disease or adult respiratory distress syndrome), obesity, or inflammatory bowel syndrome. The nucleic acids are also useful for identifying nucleic acids can also be used as primers and templates for the recombinant production of disorder-associated peptides or polypeptides, for chromosome and gene mapping, or for tissue distribution studies. The present sequence represents a gene 216 specific PCR primer used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                       scope of the invention
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Gaps
                               ;
0
   Score 13.8; DB 1; Length 20;
Pred. No. 8.6e+02;
0; Mismatches 2; Indels
    0.8%;
Query Match
Best Local Similarity 88.2
Wordhes 15; Conservative
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AAD55476 standard; DNA; 20 AAD55476; RESULT :

BP

(first entry) 07-AUG-2003 Human FGFR-3 antisense oligonucleotide, ISIS #125180.

Human, antisense, fibroblast growth factor receptor 3; prophylaxis; developmental disorder; hyperproliferative disorder, antisense therapy; FGFR-3; ACH; JTK4; CEK2; cancer; phosphorothioate; ss.

Homo sapiens Synthetic.

/mod_base= OTHER /note= "Phosphorothioate backbone; All cytidine residues are 5-methylcytidines" Location/Qualifiers ď 1. .20 /*tag= Key modified_base

-methoxyethyl (2'-MOE) nucleotides" /mod_base= OTHER /note= "2'-methoxyethyl (2'-MOE) 16. .20 mod_base= OTHER note= "2 -methor υ *tag= b '*tag= modified_base modified base

nucleotides"

WO2003023004-A2

20-MAR-2003.

06-SEP-2002; 2002WO-US028549

10-SEP-2001; 2001US-00953047.

(ISIS-) ISIS PHARM INC

Monia BP, Wyatt JR;

WPI; 2003-313244/30.

Novel compound targeted to a nucleic acid molecule encoding fibroblast

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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding fibroblast growth factor (FGF) receptor 3 (also known ser PGFR-3, ACH, JUYK4 and CEK2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with FGFR-3 such as developmental disorders or by-perpoilerative disorders, especially cancer of colorectal, bladder, bone, lung, cervical, breast or skin. They are useful as research reagents, therapeutics, prophylaxis, kits and diagnostics, and as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion of the genes expressed within cells and tissues. They are also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human FGFR-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
growth factor receptor 3, useful for inhibiting the expression of receptor and for treating an animal having cancer or developmental disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 8 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                             Example 15; Page 79; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 88.2
les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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RESULT 1123 ACF57208/c ID ACF57208 standard; DNA; 20 BP. ACF57208;

977 GAGACCTCAAGCCCCAG 993

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2 GAGACCCCAAGCCCCTG

Human LAMA3 reverse PCR primer SEQ ID NO:8. 16-OCT-2003

Human, mouse, skin structure, skin, laminin 5 chain gene, LAWA3; LAMB3, LAMC2, extracellular matrix component, matrix metalloproteinase, MMP-1, MMP-2; MMP-3; MMP-9; TIMP-1, TIMP-2; TIMP-3; collagen, PCR primer, ss.

Homo sapiens Synthetic. JP2002330792-A.

19-NOV-2002

15-JAN-2002; 2002JP-00006797

15-JAN-2001; 2001JP-0006952.

(SHIS) SHISEIDO CO

WPI; 2003-407328/39.

A method and a kit for determination of expression of mRNA or cDNA of protein participating in the maintenance of skin structure.

Claim 1; Page 2; 34pp; Japanese.

The present invention describes a method and a kit for determining the expression of mRNA or cDNA of a protein participating in the maintenance of skin structure. The method is quantitative, simple and accurate in the determination of extracellular matrix components of laminin 5 chain genes LAMA3, LAMB3 and LAWC2, matrix metalloproteinases MMP-1, MMP-2, MMP-3 and MMP-9, VII collagen, type I collagen alpha 1 chain, type IV collagen alpha 1 chain, ACF57290 represent PCR primers and probes used in the method of the

invention

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RESULT 1124

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Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating non-allergic inflammatory diseases, such as psoriasis, ecze allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                             Immunostimulatory nucleic acid #972.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 35; 229pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunostimulatory nucleic acid #929.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1547 GCCTTCGGTCTTCGTCG 1563
                                         BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-2001; 2001US-0279642P
                                       ACH03337 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB37315 standard; DNA; 20
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                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 15; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (KRIE/) KRIEG A M. (BERG/) BERG D J.
                                                                                                                                                                                                                                                                                                                                       JS2003050268-A1.
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                                                                                                                     25-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                             13-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Krieg AM,
                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is that of an antisense primer for Fas-associated protein with death domain (FADD). Use with the sense primer given in ACPOSTAS generates a 695 bp product. Semi-quantitative RT-PCR was used to determine levels of FADD RNA in thyroids of gen transgenic mice during various stages of tumour development. These mice provide models of human. Approved tumours. FADD expression was shown to decrease, in some cases to zero, during tumour development. It therefore provides a marker for the absence of in vivo tumour. FADD proteins are secreted from tumour cells. A low cellular amount and a high extracellular amount of FADD proteins are prognostic of resistance to chemcherapy. The invention provides methods for determining a status of tumour absence/presence, and for prognosis of the resistance of a tumour to chemotherapy on the basis of
                                                                                                                                                                                                                                                                                                                                                                                                              FADD; mouse; tumour; marker; diagnosis; prognosis; thyroid; PCR; primer;
                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Use of Fas associated protein with dead domain, and cellular phosphorylated p38-mitogen activated protein kinases as a biological indicator of tumor status.
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                                 Score 13.8; DB 1; Length 20;
Pred, No. 8.6e+02;
0; Mismatches .2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred, No. 8.6e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chiocchia G, Tourneur L, Feunteun J, Michiels F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 7 A; 7 C; 4 G; 2 T; 0 U; 0 Other;
Sequence 20 BP; 9 A; 7 C; 4 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (INRM ) INSERM INST NAT SANTE & RECH MEDICALE (CNRS ) CNRS CENT NAT RECH SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 50; 118pp; English.
                                                                                                                 1113 TGACATCCTGCTTGGGT 1129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1656 CCACACCCCTCACAGGG 1672
                                                                                                                                                                                                                                                         87
                                   0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-DEC-2002; 2002WO-EP014906.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-DEC-2001; 2001EP-00403359.
22-OCT-2002; 2002EP-00292619.
          Query Match
Best Local Similarity 85...
                                                                                                                                                     20 reretrecreerieger
                                                                                                                                                                                                                                                     ACF05737 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                         FADD antisense PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-645962/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003056340-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          these findings
                                                                                                                                                                                                                                                                                                                                   06-NOV-2003
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                                                                                                                                                                                                                                                                                             ACF05737;
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The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                                     0.8%; Score 13.8; DB 1; Length 20;
llarity 88.2%; Pred. No. 8.6e+02;
Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
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ccacagrecreacage 20

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The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
/note= "Phosphorothioate backbone and all cytosines are
-methyl cytosines"
                                                                                                                                                                                                                                          Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antinsense oligonucleotide targeting mouse C3 component, ISIS140104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   septic shock; multiple organ failure; hyperacute organ failure;
autoimmune disorder; CNS inflammation; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse; 88; antisense; complement component C3; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= a
/mcd_base= OTHER
/note= "2'-methoxyethyl nucleotides"
/f6 . .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "2'-methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                            an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                     Bratzler RL, Petersen DM, Fouron Y;
                                                                                                                                                                                                                                                                                Disclosure; Page 19; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1547 GCCTTCGGTCTTCGTCG 1563
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                                                                            02-FEB-2001; 2001US-00776479
                                                                                                       03-FEB-2000; 2000US-0179991P
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                                                                                                                                 BRATZLER R L.
PETERSEN D M.
FOURON Y.
                                                                                                                                                                                                               WPI; 2003-657977/62.
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Matches 15; Conserv
                         US2003087848-A1
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The invention relates to a compound 8-50 mucleobases in length targeted to a nucleic acid molecule encoding complement component C3. The compound of specifically hybridises with the nucleic acid molecule encoding complement component C3 and inhibits the expression of complement component C3 and inhibits the expression of complement component C3. Also included are a composition comprising the expression of a pharmaceutical carrier or diluent, inhibiting the expression of complement compound cited above) and treating an animal camplement component C3 in cells or tissues (complement component C3 in cells or tissues (complement component C3 complement component C3 in complement component C3 in complement component C3 in complement component C3 is inhibiting the expression of complement component C3 is inhibiting the expression of complement component C3 is inhibiting an animal having a disease or condition associated with complement component C3 such as an autoimmune disorder (e.g. multiple sclerosis), an infection, or atherosclerosis, inflammation, septic shock, multiple organ failure, or useful as research reagents and diagnostics, in distinguishing functions continue members of a biological pathway, or for preventing or delaying intervance organ failure and complements of a various members of a biological pathway, or for preventing or delaying an entire component C3 in the present sequence is an entired and entire component C3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense; ss; human; eukaryotic translation initiation factor 2C 1; EIF2C1; Co-EIF2C; Golgi ER protein 95kDa; GERp95; Q99; gene therapy; hyperproliferative disorder; familial hypercholesterolaemia; cancer; polycystic kidney disease; cystic fibrosis; progeriod syndrome; cytostatic; antilipaemic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides targeted to a nucleic acid molecule encoding complement component C3, useful for treating a disease or condition associated with complement component C3, e.g. autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
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                                                                                                                                             23-OCT-2001; 2001US-00001076.
                                                                                                                                                                                                                    23-OCT-2001; 2001US-00001076.
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                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
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US2003096775-A1.
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                                                                      22-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                     This invention relates to novel antisense oligonucleotides that modulate the expression of human eukaryotic translation initiation factor 2C 1 (EIF2C1). EIF2C1 is located on chromosome 1934-35, and is also known as Co-eIF2C, eIF2C1 Golgi Ex protein 95kDa, GER995 and 199. It is an intracellular membrane associated protein thought to be involved in cellular differentiation, such that altered expression of EIF2C1 can affect cell growth, morphology and tumourigenicity. Accordingly, antisense oligonucleotides that inhibit the expression of EIF2C1 in cells including hyperproliferative disorders, damilial hypercholesterolaemia and cancer, as well as polycystic kidney disease, cystic fibrosis and progeriod syndrome. As such, the oligos of the present invention can be described as having cytostatic and antilipaemic activities. This oligonucleotide sequence is an antisense oligo used to inhibit expression of the human eukaryotic translation initiation factor 2C 1 (EIF2C1) DNA
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                                                                                                                                                                                                                                                                                    New compound, having a sequence targeted to a nucleic acid encoding human collapsin response mediator protein 2, useful for preparing a composition for treating hypercholesterolemia or hyperproliferative disorder, e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, ss; PCR; retinal pigment epithelial-derived neurotrophic factor; PEDF; tumour, ocular disease; neuronal cell pathology; serine protease; blood coagulation; thrombosis; bacterial infection; parasitic infection; elastosis; vascular disorder; fibrinoid formation; coagulation disorder; arteriosclerosis; ischaemia; arthroses diabetes; emphysema; arthritis;
                                               /mod_base= OTHER
/note= "OTHER= phosphorothioate backbone, where 1-5 and
16-20 are 2' methoxyethyl nucleotides. All cytidines are
5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human retinal pigment epithelial-derived factor (PEDF) PCR primer #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 0.8%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 8.6e+02; es 15; Conservative 0; Mismatches 2; Indels
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             Location/Qualifiers
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                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
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of the invention.
                                                                                                              WO2003040321-A2
              Key
modified_base
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Best Local (
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The invention relates to a composition comprising purified retinal comprisions comprise ADB99089, ADB99080 requested neurotrophic factor (PEDF). The PEDF controlled are professed comprises ADB99089, Human PEDF is encoded by ADB99088 Also included are purifying PEDF, producing PEDF comprising expressing the DNA sequence concomplinate DNA molecule comprising a comprising a PEDF in a host cell, a recombinant DNA molecule comprising a peope, a recombinant DNA molecule comprising a retinal PEDF CDNA, a host cell containing the vector, a recombinant DNA molecule comprising a retinal PEDF CDNA, a host cell containing the vector, a recombinantly produced PEDF protein siolated or is free from the risks normally associated with proteins isolated or continued a naturally occurring source organism and a purified human cell containing the vector, colar disease, neuronal cell pathologies, or conditions resulting from the activity of serine proteases, escing parasitic infection, colared proteins isolated retinal infection, parasitic infection, alastosis, vascular disorders involving fibrinoid formation, coagulation disorders, arteriosclerosis, isohemmia, arthrise septic shock, lung diseases, excessive complement activation, ulcers, ulcerative colitis, pancreatitis, fibrinolytic disease, arthropathy, bone resorption, proteases, the present sequence is a PCR primer used to isolate genomic DNA encoding human retinal pigmented epithelium derived repitment used to isolate genomic CDNA encoding human retinal pigmented epithelium derived repitment used to isolate genomic CDNA encoding human retinal pigmented epithelium derived repitment used to isolate genomic CDNA encoding human retinal pigmented epithelium derived repitment used to isolate genomic CDNA encoding human retinal pigmented epithelium derived repitmented epithelium derived preciped to the preciped to the preciped to the preciped to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New purified retinal pigmented epithelium derived neurotrophic factor composition, useful for treating tumors, i.e. retinal tumor, ocular disease, neuronal cell pathologies, coagulation disorders or arteriosclerosis.
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septic shock, lung disease; complement activation; ulcer; ulcerative colitis; pancreatitis; psoriasis; fibrinolytic disease; arthropathy; bone resorption; hypertension; congestive heart failure; cirrhosis; protease allergy; chromosome 17p13.1-pter; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tombran-Tink J, Steele FR, Chader GJ, Becerra SP, Johnson LV;
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88.2%; Pred. No. 8.6e+02;
Attive 0; Mismatches 2; Indels
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92US-00952796.
95US-00520373.
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Best Local Similarity 88.2
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (TOMB/) TOMBRAN-TINK J. (STEE/) STEELE F R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (JOHN/) JOHNSON L V.
(RODR/) RODRIGUEZ I R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STEELE F R.
CHADER G J.
BECERRA S P.
JOHNSON L V.
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24-SEP-1992;
29-AUG-1995;
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                                                                                                                                                                                                                             Homo sapiens
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(BECE/)
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The invention comprises antisense oligonucleotides that are targeted to the nucleic acid encoding transforming growth factor beta (TGF-beta) receptor II. The antisense oligonucleotides of the invention are useful for treating: hyperproliferative disorders (e.g. breast cancer), or an autoimmune disorder (e.g. rheumatoid arthritis). The present DNA sequence represents a 2'-C-methoxyethyl gapmer oligonucleotide with a phosphorothioate backbone that is targeted to human TGF-beta receptor II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New compound having a sequence targeted to a nucleic acid encoding Transforming growth factor beta-receptor II, useful for preparing a composition for trating hyperproliferative disorder e.g., lung, liver, colon or gastric cancer.
                                                                                                                                                 human; antisense oligonuclectide;
transforming growth factor beta receptor II; TGF-beta receptor II;
typerproliferative disorder; breast cancer; autoimmune disorder;
rheumatoid arthritis; 2.0-methoxyethyl gapmer;
phosphorothioate backbone; ss.
                                                                                                                 Human TGF-beta receptor II targeted antisense oligonucleotide #52
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88.2%; Pred. No. 8.6e+02;
iive 0; Mismatches 2; Indels
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775/c
ADC65775 standard; DNA; 20 BP.
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                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murray SF, Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-175279/17.
                                                                                                                                                                                                                                                                                           WO2003000656-A2
                                                                                                                                                                                                                                                         Homo sapiens.
                                                                              18-DEC-2003
                                            ADC65775;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Loca
Matches
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proteins (II) from Lollum perenne and Feetuca arundinacea which are active in lighth, fructan and tannih blosynthetic pathways. Also described (1) an isolated oligonucleotide probe or primer comprising at least 10 contiguous residues complementary to 10 contiguous residues of (I); (2) a kit comprising the oligonucleotide probe or primer; (3) a genetic construct comprising (I); (4) a transgenic plant cell comprising the genetic construct of (3); (5) a plant or its seed, fruit or progeny comprising the transgenic plant cell of (4); (6) modulating one or more of the lighin, fructan or tannih compositions; and (8) modifying the activity of (II) involved in a lighin, fructan or tannih compositions; tannih blosynthetic pathway in a plant. (1) can be used for modulating the blosynthesis of lighin, fructan or tannih in a plant. The present sequence is used in the exemplification of the present invention.

New polynucleotide encoding polypeptides from Lolium perenne or Festuca arundinacea, useful for modulating the biosynthesis of lignin, fructan or tannin in a plant.

Glenn

Norriss MG,

Shenk MA,

Gibson JB,

Demmer J, Forster RL, Saulsbury KM, Hall C; WPI; 2003-441544/41.

(GENE-) GENESIS RES & DEV CORP LTD. (WRIG-) WRIGHTSON SEEDS LTD.

07-NOV-2002; 2002WO-NZ000239 07-NOV-2001; 2001US-0337703P.

Lolium perenne. Schedonorus arundinaceus.

WO2003040306-A2.

L5-MAY-2003

present invention describes isolated polynucleotides (I) encoding

Example 8; SEQ ID NO 217; 240pp; English.

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 0.8%; Score 13.8; DB 1; Length 20;
18.2%; Pred. No. 8.6e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                Yeast CYP52A5A/B genes 5' region RT-PCR primer #1.
                                                                     851 TGGACAAGGACCTGAAG 867
                                                                                                                                                                                             BP
                                                                                                                                                                                             ADC45046 standard; DNA; 20
                                                                                                                                                                                                                                                                 (first entry)
Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                 18-DEC-2003
                                                                                                                                                                                                                              ADC45046;
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Lolium perenne; Festuca arundinacea; lignin; fructan; tannin; biosynthetic pathway; plant; PCR primer; ss.

Synthetic

Tannin biosynthesis gene related PCR primer SEQ ID NO:217.

(first entry)

18-DEC-2003

SXXXXXXXXXXXX

ADC68507;

Bb

ADC68507 standard; DNA; 20

RESULT 1131

ADC68507

19 ccarcirircrococicc 3

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Candida tropicalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2003049821-A1.
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New nucleic acid encoding cytochrome P450 and NADPH reductase enzymes (e.g. CPRA, CPRB or CYP52A1A), useful for producing dicarboxylic acids that may be utilized as industrial intermediates in manufacturing
                                                                                                                                                                                                                                                                                 Example 11; SEQ ID NO 47; 196pp; English
                                        01-MAX-1998; 98US-0083798P.
05-OCT-1998; 98US-0103099P.
10-MAX-1999; 99US-012355FP.
30-APR-1999; 99US-0030620.
12-OCT-2001; 2001US-00976800.
                         03-MAY-2002; 2002US-00138838
                                                                                                                                                                                                                                              (e.g. CPRA, CPRB or CY)
that may be utilized a
diesters and polymers.
                                                                                            (WILS/) WILSON C R.
(CREAT D L.
(ERRI/) ETRICH L D.
(ESHO/) ESHOO M.
(MADD) MADDURI K M.
(CORN/) CONNETT C A.
(BREN/) BRENNER A A.
(TAMG) TAMG M.
(LOPE), LOPER J C.
(GLEE/) GLEER J C.
                                                                                                                                                                                                                      WPI; 2003-777150/73.
                                                                                                                                                                                                    Brenner AA,
                                                                                                                                                                                             Wilson CR.
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The invention relates to an isolated mucleic acid selected encoding candida tropicalis omega oxygenase complex enzymes (Cytochrome P450 monooxygenase (CYP) and MADPH reductase enzymes (Cytochrome P450 monooxygenase (CYP) and MADPH reductase enzymes (CYP) designated CPRA, CPPS2A3A, CYPS2A3A, CYPS2A3A, CYPS2A3A, CYPS2A3A, CYPS2A3A, CYPS2A3A, CYPS2A3A, CYPS2A3A, CYPS2A3A, CYPS2A3B, CYPS2A3A, CYPS2A3B, CYPS2A3A, CYPS2A3B, CYPSA3B, COMPETIENCE OF COMPETIENCE COMPETIENCE COMPETIENCE CONTCOL POX MANA in response to exogenously added control Competience of expense control POX mana in response to exogenously added control Competience control Competienc Gaps ô Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;

Query Match

0.8%; Score 13.8; DB 1; Length 20;
Best Local Similarity 88.2%; Pred. No. 8.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels

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1010 AGAGGGGAGAGCTCAAG 1026

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ADC45616 standard; DNA; 20 ADC45616; RESULT 1133 ADC45616 SASSES

(first entry) 18-DEC-2003

PCR; Primer; 88; yeast; omega oxygenase complex; cytochrome P450 monoxygenase; CPP; NADPH reductase enzymes; CPR; CPRA; CPRS; CPRS; CPRS; CPRS; CYPS2A1A; CYPS2A1A; CYPS2A3B; CYPS2A5A; CYPS2A5B; CYPS2A5A; CYPS2A5B; CYPSZA5B; CY feast CYP52A5A/B genes 5' region RT-PCR primer #1

Candida tropicalis.

US2003049822-A1.

13-MAR-2003

03-MAY-2002; 2002US-00139031

01-MAY-1998; 98US-0083798P. 05-OCT-1998; 98US-0103099P. 10-MAR-1999; 99US-0123555P. 30-APR-1999; 99US-00302620. 12-OCT-2001; 2001US-00976800.

WILSON C R.

(MILS/)

Madduri KM, Cornett CA,

Eirich LD, Eshoo M, Loper JC, Gleeson M;

Loper JC,

Craft DL, , Tang M, 1

MADDUEL K M.
CORNETT C A.
BRENNER A A.
TANG M.
LOPER J C.
GLEESON M. CRAFT D L. EIRICH L D. ESHOO M. (CRAF/)
(BIRI/)
(BSHO/)
(MADD/)
(CORN/)
(BREN/)
(TANG/)
(LOPE/)

Madduri KM, Eirich LD, Eshoo M, Loper JC, Gleeson M;

Cornett CA;

Craft DL, Tang M, Wilson CR, Brenner AA,

WPI; 2003-765370/72.

New nucleic acid encoding cytochrome P450 and NADPH reductase enzymes (e.g. CPRA, CPRB or CYP52A1A), useful for producing dicarboxylic acids that may be utilized as industrial intermediates in manufacturing diesters and polymers.

Example 11; SEQ ID NO 47; 196pp; English.

The invention relates to an isolated nucleic acid selected encoding candida tropicalis omega oxygenase complex enzymes (Cytochrome P450 monooxygenase (CYPS and NADPH reductase enzymes (CPR) designated CPRA, character (CPR) and NADPH reductase enzymes (CPR) designated CPRA, CPPS2A5B, CYPS2A5B, CY

Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;

ö Gaps ö 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; tive 0; Mismatches 2; Indels Best Local Similarity 88.2 Matches 15, Conservative Query Match

1010 AGAGGGGAGAGCTCAAG 1026

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"mod_base= OTHER
note= "Phosphorothioate backbone and all cytidines are 5
.methyl cytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel compound hybridizing with nucleic acid molecule encoding CD81 and inhibiting the expression of CD81, useful for treating infections and disease associated with expression of CD81 such as inflammation disorder.
                                                                                                                                              Antisense; ss; human; CD81; TAPA-1; tetraspanin; viral infection; cocaine addiction; autolmmune disorder; antinflammatory; antibacterial; virucide; antiparasitic; inflammatory disorder; parasitic infection; bacterial infection.
                                                                                                                                                                                                                                                                                                                                              /mod_base= ОТНЕR
/note= "2'-methoxyethyl nucleotide"
                                                                                                                                                                                                                                                                                    *tag= a
*tag= a
mod_base= OTHER
/note= "2'-methoxyethyl nucleotide"
|6. .20
|*tag= c
                                                                                                                               Human CD81/TAPA-1 antisense oligonucleotide #60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; SEQ ID NO 72; 55pp; English.
                                                                                                                                                                                                                      Location/Qualifiers
1. .20
                   AGAGGCAGGCTCAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                        10-DEC-2001; 2001US-0006430
                                                            .600/c
ADC35600 standard; DNA; 20
                                                                                                            18-DEC-2003 (first entry)
                                                                                                                                                                                                                                            *tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dobie K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-810907/76.
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                                                                                                                                                                                                                        Key
modified_base
                                                                                                                                                                                                                                                                                     modified_base
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                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                 19-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Graham MJ,
                                                                                        ADC35600;
                                                 RESULT 1134
                                                           ADC35600
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The invention relates to a compound (antisense oligomuclectide) hybridising with the eighth nucleobase portion of an active site on a nucleic acid molecule encoding CDB1 (also known as TMpA-1, a tetraspanin) and inhibiting the expression of CDB1. Also included is a composition comprising the antisense oligomuclectide and a carrier or a diluent. The antisense oligomuclectide in useful for inhibiting the expression of CDB1 in cells or tissues. The antisense oligomuclectide is useful for intracting infections preferably viral, bacterial and parasitic and disease or condition is characterised by chemical dependency (e.g. cocaine addiction). The present sequence is a CDB1 antisense oligonuclectide of the invention.

Sequence 20 BP; 6 A; 4 C; 8 G; 2 T; 0 U; 0 Other;

Query Match

DB 1; Length 20; 0.8%; Score 13.8;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention comprises oligonucleotides for detecting and identifying subtypes of human papilloma virus (HPV) contained in a sample. The oligonucleotides of the invention are useful for simultaneously detecting and identifying subtypes of HPVs. The present DNA sequence represents an HPV detection oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detector for identifying human papilloma virus subtypes, comprises carrier having two parts carrying first and second oligonucleotides that respectively hybridize with DNA contained in first and second subtypes of the virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88
                                                                                                                                                                                                          Human papillomavirus type 6 (HPV 6) detection oligonucleotide #2.
                                                                                                                                                                                                                                 probe; human papilloma virus; HPV; detection; identification; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        probe; human papilloma virus; HPV; detection; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human papillomavirus type 6 (HPV 6) detection oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                   Fan
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0.8%; Score 13.8; DB 1; Length 20;
Best Local Similarity 88.2%; Pred. No. 8.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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88.2%; Pred. No. 8.6e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   Lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 6 A; 7 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                   н, <sub>С</sub>
                                                                                                                                                                                                                                                                                                                                                                                                                    Chan
                                                                                                                                                                                                                                                                                                                                                                                                                 Lin R, You C, Huang H, Lee B, Shih C, Yeh C, Kao Y, Pan C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID NO 466; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                          (KING-) KING CAR FOOD IND CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19
                                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                   10-OCT-2001; 2001EP-00123379.
                                                                                                                                                                                                                                                                                                                                          10-OCT-2001; 2001EP-00123379.
                                                                                                                                                                                                                                                           Human papillomavirus type 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC84235 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 ccgraacracarcricc
                                         1700 ACTCTCTGCCTACCTGC
                                                       ADC84236 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JAN-2004 (first entry)
                                                                                                                                                                                    entry)
  Best Local Similarity 88.23
Matches 15; Conservative
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                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                        EP1302550-A1.
                                                                                                                                                                                    01-JAN-2004
                                                                                                                                                                                                                                                                                                                16-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC84235;
                                                                                                                                                          ADC84236;
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Heu H,
                                                                                                                                    a
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Example 3; SEQ ID NO 35; 308pp; Japanese.

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Fan C;

Lin Y,

Lee H, Chan P;

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Human papillomavirus type

EP1302550-A1

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The invention comprises oligonucleotides for detecting and identifying subtypes of human papilloma virus (HPV) contained in a sample. The oligonucleotides of the invention are useful for simultaneously detecting and identifying subtypes of HPVs. The present DNA sequence represents an HPV detection oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     angiogenesis inhibitor; cytostatic; antiinflammatory; cancer; coarian disease; diabetic retinopathy; inflammatory; ZAQ; Bv8; ISE; ss; PCR; primer; RBv8-WR2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detector for identifying human papilloma virus subtypes, comprises carrier having two parts carrying first and second oligonucleotides tha respectively hybridize with DNA contained in first and second subtypes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Match 0.8%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Fred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Angiogenesis inhibitor-related PCR primer RBv8-WR2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 5 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                     Pan C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 465; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                               ree
                                                                                                                                                                                                                                                                                                                                                     Huang H,
                                                                                                                                                                                                                                                                                                   (KING-) KING CAR FOOD IND CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Takatsu Y;
                                                                                                                                                                                                                                                                                                                                                                                           Kao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1677 CCCCAACTACATCTTCC 1693
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                                                                                                                                                                             10-OCT-2001; 2001EP-00123379.
                                                                                                                                                                                                                                      10-OCT-2001; 2001EP-00123379
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                                                                                                                                                                                                                                                                                                                                                     You C, H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-432398/41.
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                                                                                                                                                                                                                                                                                                                                                        Lin R, )
Shih C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003066860-A1.
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                                                                                                                 16-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                        Lin C,
Hsu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1137
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Gaps

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2; Indels

cancer, ovarian

Angiogenesis inhibitors for treatment and prevention of diseases and inflammatory disease.

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The invention relates to a novel angiogenesis inhibitor comprising a compound that inhibits the activity of an amino acid sequence given in the specification. Angiogenesis-related proteins Bv8, ZAQ and 15E were utilised within the method of the invention. The molecules of the invention demonstrate cytostatic and antiinflammatory activities whilst the method may be useful for treatment and prevention of cancer, ovarian diseases, diabetic retinopathy and inflammatory disease. The current sequence is that of the angiogenesis inhibitor-related PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Investigating male genetic infertility, useful for diagnosis e.g. for assessing sultabilty for in vitro fertilization, based on multifactorial analysis of infertility-related mutations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for investigating genetic infertility or predisposition in males. The method involves selecting at least two infertility-associated mutations which are recessive or intermediate that are associated with infertility in the heterozygous state and/or only in the homozygous state. Preferably at least one azoospermia factor is detected which may be lost by microdeletions intervals 5 or 6 of the Y-chromosome. Also any of several hundred mutations, listed, present in the cystic fibrosis transmembrane conductance regulator (CTFR), Kallmann syndrome (KAL1), androgen resistance (AR) or steroid 21-hydroxylase (CYP21) genes may be detected.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer, male infertility; infertility-associated mutation; azoospermia factor; Y-chromosome; cystic fibrosis transmembrane conductance regulator; CTFR; Kallimann syndrome; KALI; androgen resistance; steroid 21-hydroxylase; CYP21; microarray; quantitative trait locus; in vitro fertilization; oligospermia; ss.
                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 8.6e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 7 C; 3 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 13; SEQ ID NO 73; 110pp; German.
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                                                                                                                                                                                                                                                                                                                                                      CTGAAGCAGTACCTGGA
                                                                                                                                                                                                                                                                                                                                                                                          creasecadeacreda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD42212 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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Best Local Similarity 88.2'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cullen P, Seedorf U;
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                                                                                                                                                                                                                                                                                                                                                      862
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ADD42212
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Sequence 20 BP; 6 A; 3 C; 7 G; 4 T; 0 U; 0 Other;

Gaps . 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 8.6e+02; tive 0; Mismatches 2; Indels Query Match Best Local Similarity 88.2⁵ Matches 15; Conservative

577 GTCAGCCTATCTGAGAT 593 GGCAGCCTATGTGAGAT 20

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Example C; SEQ ID NO 318; 447pp; English.

asthma, or infections.

RESULT 11 ADE28941/

ADE28941;

ADE28941 standard; DNA; 20 BP

(first entry) 29-JAN-2004

Reverse Ag2597 RT-PCR primer used to amplify human NOV RNA.

NOVX; antidiabetic; anorectic; cardiant; hypotensive; antidiabetic; anorectic; cardiant; hypotensive; antidateriosclerotic; virucide; antibacterial; fungicide; protozoacide; notoropic; neuroprotective; antibarkinonian; anticonvulsant; osteopathic; antidiatiammatory; dermatological; antiasthmatic; antilipaemic; metabolic; diabetes; obesity; infectious; anoreaxia; cancer; cardiovascular; hypertension; atherosclerosis; neurodegenerative; Alzheimer's disease; Parkinson's; epilepsy; immune; osteoarthritis; haemopoietic; inflammatory skin; asthma; dyslipidaemia; neurogenesis; call differentiatin; proliferation; haemopoiesis; wound healing; angiogenesis; gene therapy; chromosome mapping; tissue typing; human; NOV; PCR; primer; ss; RT-PCR.

Homo sapiens.

WO2003040330-A2.

15-MAY-2003

05-NOV-2002; 2002WO-US035536

2001US-0336600P. 2001US-0338285P.

2001US-0342592P. 2001US-0344297P. 2001US-0344903P. 2002US-0373288P. 2002US-0380981P. 2001US-0341346P. 2001US-0341477P. 2001US-0341540P. 17-APR-2002; 15-MAY-2002;

Alsobrook JP, Alvarez E, Anderson DW, Baron M, Boldog FL, Bisen A; Blerman SJ, Chapoval A, Dhanabal M, Edinger SR, Eisen A; Ellerman K, Ettenberg S, Gangolli EA, Gerlach VL, Gorman L, Grosse WM, Guo X, Hackett C, Ji W, Rekuda R, Khramtsov NV; Lepley DM, Li L, Macdougall JR, Malyankar UM, Mazur A, Mcqueeney K; Mezes PS, Miller CB, Millet I, Mishra VS, Padigaru M, Patturajan M; Pena CEA, Peyman JA, Rastelli L, Rieger DK, Shenoy SG, Shimkets RA; Smithson G, Starling G, Spytck KA, Stone DJ, Tchernev VT, Twomlow N; Vernet CAM, Zerhusen BD, Zhong M; New isolated NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders, e.g. osteoarthritis, obesity, atherosclerosis, cancer, Parkinson's disease, 17-MAY-2002; 2002US-0381495P.
28-MAY-2002; 2002US-038354P.
29-MAY-2002; 2002US-0383744P.
29-MAY-2002; 2002US-0383029P.
29-MAY-2002; 2002US-0384024P.
26-AUG-2002; 2002US-0401788P.
26-AUG-2002; 2002US-0401788P.
31-OCT-2002; 2002US-040533P. (CURA-) CURAGEN CORP. WPI; 2003-441555/41. Probes for the mutated genes and/or native nucleic acid, or their complementary strands, are fixed to a carrier, particularly as a microarray, then tested for hybridization with oligomortectides from or synthesized from, a patient sample and hybridization detected.

Multifactorial analysis is by standard statistical methods, particularly the quantitative trait locus method. The method is used to diagnose inherited male infertility or predisposition to its, especially in conjunction with in vitro fertilization programs, e.g. for assessing intracytoplasmic sperm injection method. Analysis of many mutations intracytoplasmic sperm injection method. Analysis of many mutations improves diagnosis of the genetic basis of male infertility, including polygonic origins (complex interactions between different heterozogotic mutations). A chip for analyzing genetic infertility in males comprises coligomuclectides that represent known mutations (nonsense or missence, insertions, allelic variants deletions or rearrangements) in the cystic fibrosis transmembrane conductance regulator, Kallmann syndrome, androgen resistance and steroid 21-hydroxylase genes. ADD42140-ADD42633 represent coligonucleotides used in the microarray described in the method of the invention. NOTE: there are no SEQ ID's 133, 472 or 473 represented in the ö

The invention relates to a novel isolated NOVX polypeptide. The polypeptide of the invention demonstrates, antidiabetic, anorectic, cardiant, hypotensive, antitarteriosolerotic, virucide, antiparkinsonian, indicide, protozoacide, nootropic, neuroprotective, antiparkinsonian, anticonvulsant, osteopathic, antiarthritic, antiinflammatory demanded, protozoacide, nootropic, neuroprotective, antiparkinsonian, anticonvulsant, osteopathic, antiarthritic, antiinflammatory for demanded and antibodies may be useful for treating or diagnosing diseases including metabolic disorders such as diseases including hypertension and atheroscierosis, cardiovascular diseases including hypertension and atheroscierosis, neurodegenerative disorders such as Alzheimer's disease, Parkinson's disorders, inflammatory skin disorders, asthma and dyslipidaemia.

Furthermore, the nucleic acids and polypeptides may also be used to identify molecules that modulate or inhibit neurogenesis, cell differentiation and proliferation, haemopolesis, wound healing and angiogenesis, as well as in gene therapy. Finally, the nucleic acids may be used as hybridisation probes, in chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The current sequence is that of interior and proliferation and proliferation and promoses and the current sequence is that of interior. invention

Sequence 20 BP; 10 A; 3 C; 5 G; 2 T; 0 U; 0 Other;

0.8%;

Best Local Similarity 88.2 Matches 15; Conservative

Query Match

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Gaps

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Score 13.8; DB 1; Length 20; Pred. No. 8.6e+02; 0; Mismatches 2; Indels

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ADE52127 standard; DNA; 20 ADE52127; RESULT 1140

BP

(first entry) 29-JAN-2004

C. tropicalis CYP52A5A/B QC-RT-PCR primer #1

Yeast, 88; PCR; primer; NADPH reductase; CPR; cytochrome P450; CYP; omega-hydroxylase; dicarboxylic acid; QC-RT PCR; Quantitative competitive reverse transcriptase PCR.

Candida tropicalis US2003068800-A1, 03-MAY-2002; 2002US-00138905

10-APR-2003

C. tropicalis CYP52A5A/B QC-RT-PCR primer #1.

(first entry)

29-JAN-2004

ADE64291;

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The invention relates to isolated mucleic acids encoding cytochrome P450 (CYP) and NADPH reductase (CPR) enzymes of the omega-hydroxylase complex of candida tropicalis. Also included are the CYP and CRR proteins (Comprising CPRA, CYPS2A1A, CYPS2A2A, CYPS2A2B, CYPS2A3B, CYPS2AB, CYPSAB, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New cytochrome P450 and NADPH oxidoreductase, i.e. CPR and CYP, genes and proteins, useful for discriminating members of a gene family by quantifying the amount of target mRNA in a sample, or for omega-oxidation of long chain fatty acids.
Yeast; ss; PCR; primer; cytochrome P450; CVP; NADPH reductase; CPR; omega-hydroxylase complex; omega-oxidation; long chain fatty acid; QC-RT PCR; Quantitative competitive reverse transcriptase PCR.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eirich LD, Eshoo M, Loper JC, Gleeson M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 11; SEQ ID NO 47; 194pp; English.
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05-OCT-1998; 98US-0103099P.
10-MAX-1999; 99US-0123555P.
310-ARR-1999; 99US-0030520.
12-OCT-2001; 2001US-00976800.
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Tang M,
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BIRICH L D.
BSHOO M.
CORNETT C A.
BRENNER A A.
TANG M.
LOPER J C.
GLEESON M.
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                                                                                              Candida tropicalis
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(CRAF/)
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(ESHO/)
(MADD/)
(CORN/)
(EREN/)
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(LOPE/)
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ADE64291
ID ADE64291
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Matches
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Cornett CA;

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The invention relates to an isolated CPRA, CPRE, CYP52A1A, CYP52A2A,

CYP52A2B, CYP52A3A, CYP52A3B, CYP52A5A, CYP52A5B, CYP52A2B, CYP52A2B,

CYP52A4B, CYP52A4B, CYP6Atrome P450, CPR - NADPH reductase) of the

CYP52A4A protein (CYP - CYCOKIONOME P450, CPR - NADPH reductase) of the

Candida tropicalis owega-hydroxylase complex. Also included are the

nucleic aids encoding the CYP/CPR proteins (including their coding

regions), a vector comprising the nucleotide acid, a host cell

transfected or transformed with the vector, discriminating members of a

gene family by quantifying the amount of target mRNA in a sample and

increasing production of a dicarboxylic acid (comprising: providing a

host cell having a naturally occurring CPR/CPP proceed and culturing the

constant of a dicarboxylic acid (comprising). The CYP and

constant in media containing an organic substrate which upregulates the

cens, to effect increased production of dicarboxylic acid). The CYP and

cPR proteins, present in higher levels than normal is useful for

increasing production of dicarboxylic acids. The present sequence is a

cuantitative competitive reverse transcriptase PCR (QC-RT PCR) primer

used to assay the levels of CYP/CPR mRNA in RNA samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated CPRA, CPRE, CYP52A1A, CYP52A2A, CYP52A2B, CYP52A3A, CYP52A3B, CYP52A5B, CYP52A5B, CYP52A5B, CYP52A6B or CYP52D4A protein, useful for increasing production of dicarboxylic acid in cells.
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18.2%; Pred. No. 8.6e+02;
ve 0; Mismatches 2; Indels
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05-0CT-1998; 98US-0103099P.
10-MAR-1999; 99US-0123555P.
310-APR-1999; 99US-00302620.
12-0CT-2001; 2001US-00976800.
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188 15; Conservative
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Tang M,
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MADDURI K M.
CORNETT C A.
BRENNER A A.
TANG M.
LOPER J C.
GLEESON M.
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CRAFT D L.
EIRICH L D.
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(CRAF/)
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(LOPE/)
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Matches
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Gaps

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Hepatitis C virus antigen expressed as recombinant in E.coli - useful for diagnosis of hepatitis C virus infection.

(TOKU) TOKUYAMA SODA KK WPI; 1992-263663/32. Disclosure, Page 64; 66pp; Japanese.

X4X4X44X2X2X2XXXX

The sequences given in AAQ27030-77 are primers. These were used to amplify the claimed hepatitis C virus genes of the invention which could then be inserted into an E. coli vector. The polypeptides encoded by the vectors were useful as diagnostic reagents for type C hepatitis and they may be produced efficiently by recombinant methods

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Gaps

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y Match 0.8%; Score 13.8; DB 1; Length 21; Local Similarity 88.2%; Pred. No. 9e+02; hes 15; Conservative 0; Mismatches 2; Indels

Query Match

Matches

Sequence 21 BP; 1 A; 5 C; 13 G; 2 T; 0 U; 0 Other;

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coli; diagnostic; reagent; type C hepatitis; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genome position 6813-6833. See also AAQ03898-Q03949. (Updated on 25-MAR-
2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting and typing human papilloma-virus - using consensus primers in polymerase chain reaction to amplify particular genomic regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
0.8%; Score 13.8; DB 1; Length 21;
Best Local Similarity 88.2%; Pred. No. 9e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                            HPV6 typing probe (MY12) for use with L1 consensus primers.
                                                                                                                                                                                                                                       Papilloma-virus; consensus primer; PCR; probe; ss
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polymerase chain reaction; ss.
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                                        AAQ03910 standard; DNA; 21 BP.
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89US-00322550.
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                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Manos MM, Wright DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1990-116005/15.
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10-MAR-1989;
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                                                                                                                             25-MAR-2003
24-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                         22-MAR-1990
                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                  AAQ03910;
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RESULT 1142
                     AAQ0391(
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein contg. non-A non-B hepatitis antigen fragment - prepd. by culturing transformants transformed by vector contg. base sequence coding specified aminoacid sequences, used for detecting hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a primer for a DNA encoding a non-A non-B hepatitis virus (NANBH) or hepatitis C virus (HCV) antigen, useful for diagnosis or detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                      non-A non-B hepatitis virus; NANBH; hepatitis C virus; HCV; antigen;
diagnosis; detection; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%; Score 13.8; DB 1; Length 21; Best Local Similarity 88.2%; Pred. No. 9e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 1 A; 5 C; 13 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                            Primer for hepatitis C virus antigen DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 8; 53pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         228 GAGTGGTGGTGGTGGCG 244
228 GAGTGGTGGTGGCG 244
                                                                                             AAV05593 standard; DNA; 21 BP.
                       s chicochicochicoch
                                                                                                                                                                                                                                                                                                                                            91JP-00354708.
                                                                                                                                                                                                                                                                                                                                                                       90JP-00412020
                                                                                                                                                  22-MAY-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                (SHIM/) SHIMOTONO K.
(GREC ) GREEN CROSS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1993-260858/33
                                                                                                                                                                                                                                               Synthetic.
Hepatitis virus.
                                                                                                                                                                                                                                                                                                                                            18-DEC-1991;
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Gaps ö

> 90JP-00304417 90JP-00304417

26-JUN-1992

11-NOV-1990;

RESULT 1145

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AAQ5638:

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New hypoglycosylated recombinant glucose oxidase - produced by expressing Aspergillus GOD gene in yeast mutant with N-glycosylation defect.
                                                                                                                                                                                                                                                                                                                                      The sequence is that of a primer which was used in the construction of a plasmid for the secretion of Aspergillus niger glucose oxidase (GOD) in Saccharomyces cerevisiae. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequence is that of a PCR primer which was used in the construction of plasmid YEpL/GOD-(His)4 as part of the construction of plasmids for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amplification; secretion; plasmid; construction; glucose oxidase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yeast mutants with N-glycosylation defects - for prodn. of hypoglycosylated proteins, including recombinant proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match 0.8%; Score 13.8; DB 1; Length 21; Best Local Similarity 88.2%; Pred. No. 9e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 7 A; 10 C; 3 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmid YEpL/GOD-(His)4 construction primer.
                                                                                                                                                     (BOEF ) BOEHRINGER MANNHEIM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BOBF ) BOEHRINGER MANNHEIM GMBH. (HOFF ) ROCHE DIAGNOSTICS GMBH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 11; 31pp; German.
                                                                                                                                                                                                                                                                                                          Example 1; Page 7; 27pp; German.
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                                                                                        93DE-04301904.
                                                                                                                         92DE-04226095.
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93DE-04301932.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                                                                                       Kopetzki E, Lehnert K;
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                                                                                                                                                                                                                       WPI; 1994-049996/07.
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22-JUL-1994
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25-JAN-1993;
                        DE4301904-A1
                                                                                        25-JAN-1993;
                                                                                                                       07-AUG-1992;
                                                      10-FEB-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence is that of HPV6 typing probe MY12 for use with L1 consensus primers as part of a simple and rapid assay method for detecting and typing HPV in biological samples. (Updated on 25-MAR-2003 to correct PF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detection of genital human papilloma virus - by PCR amplification using defined consensus primer pairs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GOD; enzyme; recombinant; hypoglycosylated; Aspergillus niger; yeast; expression; Saccharomyces cerevisiae; ss.
                                                                                                                                                                                                                                                 Human papilloma virus, amplification; polymerase chain reaction; PCR; detection; assay; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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0.8%; Score 13.8; DB 1; Length 21;
Best Local Similarity 88.2%; Pred. No. 9e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wright DK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Glucose oxidase secretion plasmid construction primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 6 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Мапов ММ,
                                                                                                                                                                                                                   L1 consensus primer HPV6 typing probe MY12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wolinsky SM, Broker TR, Ting Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure, Page 8; 13pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                                                                                                                                                                                                                                                     89US-00343486.
89US-00322550.
89WO-US003747.
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(HOFF ) HOFFMANN LA ROCHE INC.
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4 ccgraacracarcricc 20
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                                                                                                  AAQ56381 standard; DNA; 21
                                                                                                                                                                    (revised)
(first entry)
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10-MAR-1989;
29-AUG-1989;
                                                                                                                                                                    25-MAR-2003
29-JUL-1994
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08-AUG-1994
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                                                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                    AAQ56381;
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RESULT 1146 AAQ56141

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The sequences given in AAT00282-394 represent DNA ligands to basic fibroblast growth factor (bFGF). These sequences were isolated using the primers and target regions given in AAG98421-29 using systematic primers and target regions given in AAG98421-29 using systematic containing a region of 30 or 40 random nucleotides flanked by constant containing a region of 30 or 40 random nucleotides flanked by constant containing a region of 30 or 40 random nucleotides flanked by constant containing a regions, were synthesized. The consent regions were designed to be amplified by the primers. The primer 3p7.1PS has 2 biotin phosphoramidites and two additional A residues covalently attached to its containing an equimolar mixture of the four nucleotides during synthesis. The random regions were created by uncleotide synthesis. Three pools of sspNNA were created that contain incleotide synthesis. Three pools of ssDNA were created that contain incleotide single stranded DNA (ssDNA) was then amplified by PCR. A selected single stranded DNA (ssDNA) was then amplified by PCR. A significant improvement in affinity of BNA liquads was seen after 10 counds of selection. Five distinct families of ssDNA were identified, based on regions of homology. Some sequences showed no obvious homology to the five families and are considered to be orphans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identification of ligands to basic fibroblast growth factor and thrombin - which can be modified for increased in vivo stability.
                                                                                                                                            Family 1; family 2; ligand; thrombin; systematic evolution of ligands by exponential enrichment; SELEX; heparin; selection; region of homology; inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.8; DB 1; Length 21; 57.1%; Pred. No. 9e+02; ative 6; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chemokine receptor X5.5 primer X5-5B (antisense).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 3 A; 4 C; 5 G; 2 T; 0 U; 7 Other;
                                                                                                           Family 2 bFGF DNA consensus ligand (experiment 3)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 21; Page 106; 236pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            84 CCGCGGCTCTGAGGTTGCTCG 104
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94US-00219012.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NEXS-) NEXSTAR PHARM INC.
                                                                 (first entry)
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Best Local Similarity 57.1
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1995-293073/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Janjic N, Gold L,
                                                                                                                                                                                                                                                                                                                                                                                                  10-FEB-1994;
28-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                          06-FEB-1995;
                                                                                                                                                                                                                                                                           WO9521853-A1
                                                                    14-AUG-1996
                                                                                                                                                                                                                                                                                                                   17-AUG-1995
                                                                                                                                                                                                                                    Synthetic.
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                            AAT00342;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The human papilloma virus (HPV) specific probes AAT10818-T10839 are used to detect, or type HPV for research or diagnostic purposes, e.g. to identify HPV that are implicated in genital or oral carcinomas. (Updated on 25-MAR-2003 to correct PF field.)
the secretion of A. niger glucose oxidase (GOD) in S. cerevisiae. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PA field.)
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                                                                                                                                                                           Gaps
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carcinomas, research, typing, HPV6, specific, MY12, ss.
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                                                                                                                             0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
tive 0; Mismatches 2; Indels
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                                                                                       Sequence 21 BP; 7 A; 10 C; 3 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; Col 15-16; 36pp; English.
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ID AAT00342 standard; DNA; 21 BP.
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89WO-US003747.
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(first entry)
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                               Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-319884/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              papilloma
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10-MAR-1989;
09-SEP-1989;
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10-APR-1996
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The invention relates to new oligonucleotide probes and primers used for the detection of human papillomaviruses (HPV) which are not genital types 6, 11, 16, 18 or 33. The probes and primers AAT4608-T44693 are esp. used to detect HPV types 26, 31, 31B, 35, 39, 40, 43, 45, 51-59 and 68 The primers can be used to detect these HPV types in conjunction with the consensus primers and typing probes AAT4733-T4906, which are based on and amplify fragments of the L1, E6, E7 and E1 regions of the HPV sequences. Detection of the amplification prodes is done with probes derived from consensus sequences found in all characterised HPV sequences. Probes AAT4762-810 are examples of HPV typing probes for identifying the amplified products generated by L1 consensus primers. This sequence is a sense probe which has specificity for HPV6 and binds to the HPV genome at position 6813. (Updated on 25-MAR-2003 to correct PF
                                           Nucleic acid hybridisation probes - specific for selected human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New oligo:nucleotide probes for human papilloma-virus - used for detecting and typing HPV and for detecting previously unknown HPV types and subtypes.
                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 21;
18.2%; Pred. No. 9e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; papillomavirus 6; HPV6; typing probe; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Greer CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human papillomavirus 6 specific typing probe MY12
                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 6 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhang TY,
                                                                                        Disclosure, Col 31-32; 96pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HOFF ) ROCHE MOLECULAR SYSTEMS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bauer HM,
                                                                                                                                                                                                                                                                                                                                                                                                                                          1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88US-00243486.
89US-00322550.
89WO-US003747.
90US-00613142.
93US-00050743.
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15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT78006 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
(first entry)
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Gravitt PB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-332084/30.
              WPI; 1996-299903/30
                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-AUG-1989;
14-NOV-1990;
20-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
07-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USS639871-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-SEP-1993
                                                             virus types
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT78006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1152
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                                                                                                                                                                                                                                                                                                                                                                               A set of internal sequencing primers (AAT35281-91) were used to sequence cDNA clone E1-C19 (see also AAT35277), which codes for chemokine receptor K5.5 (AAR99274). They were designed on the basis of previous sequencing
                                                                                                                                                                                                                                                                                    Chemokine receptor which binds MIP-1-alpha, RANTES and/or MCP-1 - useful in screening for agents to treat asthma, hay fever, eczema, allergies, atopic dermatitis, rhinitis or conjunctivits.
Chemokine receptor K5.5; MIP-1-alpha; RANTES; MCP-1; allergy; atheroma;
HIV; AIDS; graft rejection; stem cell; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhang TY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HPV typing probe MY12 for use with L1 consensus primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 21 BP; 6 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Manos MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Greer CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    754 GAAGTGTCCCTGCTCAA 770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89US-00322550.
89WO-US003747.
90US-00613142.
93US-00050743.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HOFF ) HOFFMANN LA ROCHE INC
                                                                                                                                                                                                                                                                                                                                                   Example; Fig 2; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88US-00243486
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                                                                                                                                      96WO-GB000143
                                                                                                                                                                   95GB-00001683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 GATGTGTACCTGCTCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT44762 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                              GLAX ) GLAXO GROUP LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Resnick RM,
                                                                                                                                                                                                                             Power CA;
                                                                                                                                                                                                                                                          WPI; 1996-362692/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-MAR-1989;
09-SEP-1989;
14-NOV-1990;
20-APR-1993;
                                                                                                                                                                   27-JAN-1995;
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                                                                          WO9623068-A1
                                                                                                                                      24-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-SEP-1993;
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29-JAN-1997
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                                                                                                         01-AUG-1996
                                                                                                                                                                                                                             Wells TNC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bauer HM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT44762;
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Gaps ö

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The sequence is that of a PCR primer plas which was used in the isolation of DNA encoding a major subunit of the Duffy blood group antigenic system, the gp-Fy proteins. The gp-Fy proteins are gp-Fy alpha and gp-Fy beta which are produced from the same gene via a mRNA splicing mechanism. It contains the major receptor by which Plasmodium viava enters red blood cells (RBC) and causes malaria. The proteins are thus useful in preventing malaria and in regulating RBC, renal and neural function. The protein or certain fragments of it, may also be used to generate antibodies, complementary peptides and drugs modelled on their terriary structure, useful in the same way
                               The present sequence is a human papillomavirus 6 (HPV6) specific typing probe. (Updated on 25-MAR-2003 to correct PF field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid encoding gp-Fy, Duffy antigen proteins - used to prevent vivax malaria and to regulate erythrocyte, neural or renal function.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gp-FY protein; Fyb71-81; duffy blood group; antigen; alpha; beta;
alternative splicing; RBC; red blood cell; malaria; treatment;
PCR primer; ss.
                                                                                                                                0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 21;
                                                                                                  Sequence 21 BP; 6 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence 21 BP; 6 A; 2 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.8; DB 1
88.2%; Pred. No. 9e+02;
rative 0; Mismatches
 Disclosure; Col 115-116; 94pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 17; Page 32; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NYBL-) NEW YORK BLOOD CENT INC
                                                                                                                                                                                                         1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                                               4 CCGTAACTACATCTTCC 20
                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens gp-Fy PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-US021067
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                        15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chaudhuri A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-297854/26.
                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-NOV-1996;
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                                                                                                                                                                                                                                                                                                                                                               AAV27016;
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                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                            RESULT 1153
                                                                                                                                                                    Matches
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Human papilloma probes and primers - useful for, e.g. detecting and

typing of human papilloma viruses.

Manos MM;

Bauer HM,

Greer CE,

Ting Y, Resnick RM,

WPI; 1998-192210/17

(HOFF) ROCHE MOLECULAR SYSTEMS INC.

88US-00243486. 89US-00322550. 90US-00613142. 93US-00050743.

09-SEP-1988; 10-MAR-1989; 14-NOV-1990; 20-APR-1993;

95US-00452055

26-MAY-1995;

06-JAN-1998

Human papillomavirus; HPV; HPV detection; HPV typing; L1 type-specific probe; ss.

Human papillomavirus

Synthetic

US5705627-A.

Probe MY12 for human papillomavirus typing.

(revised)
(first entry)

25-MAR-2003 04-JUN-1998

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Gaps

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AAV17380;

BP

standard; DNA; 21

AAV17380

RESULT 1154 **AAV1738**(

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents a human papillomavirus (HPV) L1 type-specific probe of the invention. This sequence may be used in conjuncture with L1 specific primers for detecting and typing HPV. Identification and typing of HPV is important as different types of HPV pose different risks for infected individuals. HPV16 and HPV18 have been more consistently identified in higher grades of cervical dysplasia and carcinoma than other HPV types. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 13.8; DB 1; Length 21;
18.2%; Pred. No. 9e+02;
.ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 21 BP; 6 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer for prostate specific antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Col 15-16; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 CCGTAACTACATCTTCC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV38524 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.2%;
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Best Local Similarity 88.2
Matches 15; Conservative
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AAV38524
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MAKAKAKA
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Gaps

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2; Indels

1146 TCAGATTGACATGTGGG 1162

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ion transport; Gitelman's syndrome; Bartter's syndrome;
hypokalaemic alkalosis; hypocalciuria; hypomagnesemia; diagnosis;
therapy; SSCP; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 51; 105pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  516 GGAGAGCTGACCCTCA 532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GGAGAAGCTGGACCTCA 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ25918 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human polymorphic region 107.
                                                                                                                                                                       97WO-US023553.
                                                                                                                                                                                                    96US-00778052
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                 Simon DB;
                                                                                                                                                                                                                                                                                             WPI; 1998-388029/33.
                                                                                                                                                                                                                                  (UYYA ) UNIV YALE.
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                                                                                                                                                                       19-DEC-1997;
                                                                                                                                                                                                      31-DEC-1996;
                                                                                                          WO9829431-A1
                                                                             Homo sapiens
                                                                                                                                        09-JJL-1998,
                                                                                                                                                                                                                                                                 Lifton RP,
                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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   ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence is a PCR primer for the gene encoding the prostate specific antigen, and were used in the method of the invention. The method is for identifying markers for a disease state, and comprises: (a) providing a clirst set of peripheral blood mRNAs from one or more normal subjects known to exhibit the disease state and a second set of peripheral blood mRNAs from one or more normal subjects to fmRNAs trom one or more normal subjects of mRNAs to provide nucleic acid amplification products; (c) comparing the sets of mRNAs to provide nucleic acid amplification products; (c) comparing the sets of an expectation products; (d) dentifying those mRNAs that are differentially expressed between normal subjects and subjects exhibiting the disease state; where a difference in quantity of expression of an mRNA is indicative of a difference in quantity of expression of an mathod of detecting a metastatic cancer disease state, any be detected, diagnosed, or a prognosts may be delivered by examining a blood sample rather than relying on a more invasive, or less state may be detected, diagnosed, or a prognosts may be delivered by examining a blood sample rather than relying on a more invasive, or less progression, status and response to therapies through monitoring of differentially expressed disease markers. The methods can be used for disease such as cancer (especially metastatic or prostate cancer), asthma. lupus erythematosus, rheumatoid arthritis, multiple sclerosis, interstitial cystitis, prostatis or other systemic or chronic conditions interstitial cystitis, prostatis or other systemic or chronic conditions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                           Identifying markers for disease states - by amplifying RNA from peripheral blood and identifying RNA which is differential expressed between normal and disease state subjects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thiazide-sensitive Na-Cl cotransporter; TSC; hTSC gene; human;
    amyloid lateral sclerosis; interstitial cystitis; prostatis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 3 A; 6 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human TSC gene exon 19 forward primer hTSCex19.
                                                                                                                                                                                                                                                                                 Veltri R;
                                                                                                                                                                                                                                                                                                                                                                                                        Example 6; Page 98; 158pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CCTCAGTCTGGGGGAGC 1477
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96US-0032701P.
97US-0041576P.
                                                                                                                                                          97WO-US022105
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                  prostate specific antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                 Ohara M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarry
                                                                                                                                                                                                                                                                                                             WPI; 1998-333350/29.
                                                                                                                                                                                                                                                    (UROC-) UROCOR INC.
                                                                                                                                                                                                                                                                                   An G,
                                                                 Homo sapiens.
                                                                                               WO9824935-A1.
                                                                                                                                                          05-DEC-1997;
                                                                                                                                                                                         06-DEC-1996;
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                                                                                                                                                                                                    12-DEC-1996;
24-MAR-1997;
                                                                                                                            11-JUN-1998
                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAV40603;
                                                                                                                                                                                                                                                                                 Ralph D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
      Ob
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Primers hTSCex19 forward and reverse (see AAV40603 and AAV40604, respectively) are designed to amplify exon 19 of the human hTSC gene (see AAV40661) that codes for thiaride-sensitive Na-Cl cotransporter TSC (see AAV20682). Both primers are located within introns of hTSC. 27 Sets of specific primers (see AAV40658) were used for SSCP analysis of hTSC. Amplified products were analysed for molecular variants by electrophorosis; and identified variants were sequenced. Complete linkage of Gitelman's syndrome with TSC was demonstrated. Identification of the molecular basis of Gitelman's syndrome allows for the genetic diagnosis of this disorder. The invention provides products and methods useful for diagnosis and treatment of Gitelman's syndrome and other ion transport
Thiazide sensitive cotransporter and ATP sensitive potassium channel genes - useful for developing products for the diagnosis and treatment of ion transport disorders, e.g. Gitelman's Syndrome or Bartter's Syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; disgnosls; prevention atherosclerotic plaque; premalignant metaplastic lesion; endomerriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .<del>.</del>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 5 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-SEP-1998,
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This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene present in a population and targets only one allelic forms of the gene present in a population and targets only one allelic forms methods can be normal somatic cells, and the first gene. The products and methods can be normal somatic cells, and the first gene. The products and methods can be caucers, atherosolerotic plaques, premalignant metaplastic or dysplastic lesions, benign tumours, endometriosis, polycystic kidney disease, and malignant cells from bone marrow transplants. AAZSSB12-ZSB825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 14; Fig 1; 605pp; English.
98WO-US005419.
                                                                                               97US-0041057P.
                                                                                                                                                                                          (VARI-) VARIAGENICS INC
                                                                                                                                                                                                                                                                                               Ledley FD,
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1998-521232/44.
                                                                                               20-MAR-1997;
19-MAR-1998;
                                                                                                                                                                                                                                                                                               Housman D,
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Stanton VP;

Sequence 21 BP; 2 A; 5 C; 10 G; 4 T; 0 U; 0 Other;

0.8%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 9e+02; atrive 0; Mismatches 2; Indels 741 CACCGCCATCCGGGAAG 757 15; Conservative Query Match Best Local Similarity Best Loca Matches ò

Gaps ;

19 CACCGCCATCCTGGGAG 3

d

AAZ30746 standard; DNA; 21 BP. AAZ30746; RESULT 1158 AAZ30746

19-JAN-2000 (first entry)

Human prostate specific antigen PCR primer #15

Prostate specific antigen, DNasel, marker; expression, diagnosis; differential, disease; cancer; metastatic; breast cancer; prostate; peripheral leukocyte; immune response; asthma; lupus erythematosus; rheumatoid arthritis; multiple sclerosis; myasthenia gravis; autoimmune thyroiditis; amyotrophic lateral sclerosis; ALS; interstitial cystitis; prostatitis; mRNA; PCR; reverse transcriptase-PCR; RT-PCR; screening; early; diagnosis; prognosis; monitoring; primer; ss.

domo sapiens Synthetic

409949083-A1

30-SEP-1999

99WO-US006488 24-MAR-1999;

Identifying markers of human disease, specifically for diagnosis of metastatic prostatic and breast cancers. An G, O'hara SM, Veltri RW; invasive methods for obtaining samples Disclosure, Page 101; 225pp; English 98US-00046894. (UROC-) UROCOR INC. 24-MAR-1998; Ralph D,

Sequence 21 BP; 3 A; 6 C; 8 G; 4 T; 0 U; 0 Other;

Gaps . 0 Query Match 0.8%; Score 13.8; DB 1; Length 21; Best Local Similarity 88.2%; Pred. No. 9e+02; Matches 15; Conservative 0; Mismatches 2; Indels

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AAX78886 standard; DNA; 21 BP RESULT 1159 AAX78886

08-SEP-1999 (first entry) AAX78886;

Human plasminogen PCR primer plg+289.

Tissue factor; human; thrombogenic; substructure; thrombose; tumour; vasculative malformation; vascular endothelium; plasminogen; PCR primer;

Homo sapiens.

WO9932143-Al.

01-JUL-1999.

3 11:01:46 2004

Mon May

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thrombogenic polypeptides used to, e.g. obliterate vasculative
                                                                         Example 8; Page 81; 97pp; English.
22-DEC-1998; 98WO-US027498
           97US-00996744
                                 Houston LL, Dickinson CD;
                                            WPI; 1999-405116/34.
                      (NUVA-) NUVAS LLC.
           23-DEC-1997;
                                                              nalformations.
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This invention describes novel thrombogenic polypeptides which comprise a thrombogenic substructure and a context-dependent entity which recognizes desired biologically susceptible sites, e.g. tumour vascular endothelium. A novel context-dependent functional entity comprises a substructure with having the ability to recognize desired biologically susceptible sites, where the entity imparts thrombogenic activity when positioned in the function-forming-context at the biologically susceptible sites, and the pritty has no thrombogenic activity absent a function-forming-context at the biologically susceptible sites, the context-dependent functional entities impart thrombogenic activity only at biologically susceptible sites. The context-dependent functional entities impart thrombogenic activity only at biologically susceptible sites. The context-dependent functional sites. They can be used to obliterate vasculative malformations or to selectively thrombose the vasculature of solid tumours. This sequence represents a human plasminogen FCR primer used in the method of the

Sequence 21 BP; 5 A; 4 C; 5 G; 7 T; 0 U; 0 Other;

ö 0.8%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 9e+02; tive 0; Mismatches 2; Indels Conservative Similarity 15; Query Match Local Matches

CTGGATGACTGTGGGAA 890 crecardactardas 20 874

Human ABC1 gene exon 7 fragment corrected sequence, SEQ ID NO:171. AAC69272 standard; DNA; 21 29-JAN-2001 AAC69272; RESULT 1160 AAC69272,

BB

(first entry)

Human ABC1 cholesterol transporter; chromosome 9931;
ATP-binding cassette; HDL deficiency disorder; high density lipoprotein;
Tangier disease; TD; familial HDL deficiency; FHA; polymorphism;
cardiovascular disease; coronary artery disease; coronary restenceis;
cerebrovascular disease; peripheral vascular disease;
Alzheimer's disease; Niemann-Fick disease; Huntington's disease;
X-linked adrenoleukodystrophy; cancer; gene therapy; genetic diagnosis;
prognosis; prophylaxis; drug screening; transgenic animal; ds.

Homo sapiens.

WO200055318-A2.

21-SEP-2000.

15-MAR-2000; 2000WO-IB000532

99US-0138048P. 99US-0139600P. 99US-0151977P. 08-JUN-1999; 17-JUN-1999; 01-SEP-1999;

(UYBR-) UNIV BRITISH COLUMBIA. (XENO-) XENON BIORESEARCH INC.

Pimstone SN; Wilson AR, Hayden MR,

WPI; 2000-587528/55.

New ABC1 polypeptide is useful for treating diseases associated with ABC1 biological activity, e.g. Alzheimer's disease, Huntington's disease and cancer.

Example; Fig 11; 229pp; English.

The invention relates to the human ABC1 cholesterol transporter protein

(B38082) and to modeld acid sequences (C69120) which encoded it. ABC1 is
a member of the ATP-binding cassette (ABC transporter) superfamily of
a member of the ATP-binding cassette (ABC transporter) representations, and plays a crucial role in cholesterol transport, particularly
intracellular cholesterol trafficking in monocytes and fibroblasts, being
involved in cholesterol trafficking in monocytes and fibroblasts, being
conversed in cholesterol trafficking in monocytes and fibroblasts,
conversed on chromosome 9q31, and mutations in this gene are associated
with two genetic PHD (High density lipoprotein) deficiency disorders,
conversed density choromary are an autosomal recessive disorder,
and in the blood correlate with a high risk of cardiovascular
disease, particularly occomary artery disease, but also cerebrovascular
disease, particularly coromary artery disease, but also cerebrovascular
disease, coromary restences of HDL has protective effects against
conversely, a high level of HDL has protective effects against
conversely, and methods of gene therapy for the treatment or prevention of
cardiovascular disease. The invention provides genetic constructs and
correlavascular disease compunishing the administration of an expression
cardiovascular disease compunishing the administration of an expression
certainvascular disease compunishing the administration of an expression
certachovascular disease, coronary are coronary artery disease,
intrafar and so the cardiovascular disease, oronary are coronary are or disease,
correbrovascular disease, coronary are appeared to trace or
correbrovascular disease, coronary restended or polymorphisms in the
cincreased risk for cardiovascular disease, oronary areary disease,
correbrovascular disease, coronary restended or polymorphisms in the
correbrovascular disease, coronary restended or polymorphisms in the fractument of disease, Human ABC1 proteins and methods or peripheral vascular
correbrovascular d human ABC1 gene exon fragments

Sequence 21 BP; 7 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

Gaps ö DB 1; Length 21; Score 13.8; DB 1; Length 2: Fred. No. 9e+02; 0; Mismatches 2; Indels 0 0.8%; Query Match Best Local Similarity 88.2 Matches 15; Conservative

ö

375 GGCTTCAGCCACGTCCT 391 Н 17 GGCTTCAGCCAGCTCCT

a

AAZ60648 standard; DNA; 21 BP. (first entry) 16-MAY-2000 AAZ60648;

RESULT 1161

PCR primer used to amplify kappa3-related opioid receptor cDNA.

Splice variant; kappa3 opioid receptor; muopioid receptor-1; KOR-3; morphine analgesia; opioid-mediated ingestive response; opioid;

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Splice variant; kappa3 opioid receptor; muopioid receptor-1; KOR-3; morphine analgesia; opioid-mediated ingestive response; opioid; manalgesic; gastrointestinal motility; respiration; immune system; endocrine system; autonomous nervous system; peristalsis regulator; body weight; neuroendocrine disorder; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer used to amplify kappa3-related opioid receptor cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
0.8%; Score 13.8; DB 1
Best Local Similarity 88.2%; Pred. No. 9e+02;
Matches 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                     681 CACAGACAACCTIGIGG 697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ60652 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                     15; Conservative
                                                                                                                                                    WPI; 2000-182421/16.
                                                 WO200004151-A2
                                                                                                   16-JUL-1998;
                                                                                                                                   Pasternak G,
                                                                  27-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ60652;
                                                                                                                                                                                      weight.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 11
AAZ60652/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ઠે
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CC Rappa3 opioid receptor (muopioid receptor-1, KOR-3). The specification describes four new axons of the KOR-3 gene, which combine to yield seven new KOR-3 splice variants of human, mouse and rat origin. These splice variants of human, mouse and rat origin. These splice variants are potential targets for modulating morphine analgesia and opioid-mediated ingestive responses. The KOR-3 polypeptide are used to opioid-mediated ingestive responses. The KOR-3 polypeptide are used to coreen compounds for opioid activity. Such compounds are potential analgesics or more generally agents that affect gastrointestinal core normality, respiration or the immune, andorime or autonomous nervous systems, e.g. regulators of parisalsis. Antagonists, agonists and ligands of KOR-3, as well as DNA vectors expressing KOR-3-encoding conclica caids, or sequences antisense to KOR-3 nucleic acids, are used to regulate morphine analgesia and body weight. The level of KOR-3 or tissue distribution of KOR-3 can be measured to diagnose KOR-3 related particularly controled disorders. Transgenic animals with extra copies of the KOR-3 gene, or with endogenous alleles deleted, are used to study loss or gain of function phenotypes New splice variants of the kappa-opioid receptor, useful in screening for selective analgesics and for regulating morphine analgesia or body Sequence 21 BP; 4 A; 7 C; 6 G; 4 T; 0 U; 0 Other; KETTERING INST CANCER RES Example 1; Page 30; 61pp; English 99WO-US015977 98US-0093002P Pan Y; WPI; 2000-182421/16. WO200004151-A2 (SLOK) SLOAN Pasternak G, 15-JUL-1999; 27-JAN-2000 Query Match weight. Best Loca Matches continuers AAZ60647-48 were used to amplify cDNA fragments of the murine kappa3 opioid receptor (muopioid receptor-1, KOR-3). The specification describes four new exons of the KOR-3 gene, which combine to yield seven new KOR-3 splice variants of human, mouse and rat origin. These splice variants are potential targets for modulating morphine analgesia and opioid-mediated ingestive responses. The KOR-3 polypeptide are used to careen compounds for opioid activity. Such compounds are potential analgesics or more generally agents that affect gastrointestinal analgesics or more generally agents that affect gastrointestinal contility, respiration or the immune, endocrine or autonomous nervous systems, e.g. regulators of peristalsis. Antagonists, agonists and ligands of KOR-3, as well as DNA vectors expressing KOR-3-encoding nucleic acids, or sequences antisense to XOR-3 nucleic acids, are used to regulate morphine analgesia and body weight. The level of KOR-3 or tissue distribution of KOR-3 can be measured to diagnose KOR-3 related pharmacological abnormalities or neurcendocrine disorders, particularly gene, or with endogenous alleles deleted, are used to study loss or gain of function phenotypes New splice variants of the kappa-opioid receptor, useful in screening for selective analgesics and for regulating morphine analgesia or body analgesic; gastrointestinal motility; respiration; immune system; endocrine system; autonomous nervous system; peristalsis regulator; body weight; neuroendocrine disorder; PCR primer; ss. DB 1; Length 21; Sequence 21 BP; 4 A; 7 C; 6 G; 4 T; 0 U; 0 Other; (SLOK) SLOAN KETTERING INST CANCER RES Example 1; Page 29; 61pp; English. 98US-0093002P Pan Y;

Human biallelic marker downstream amplification primer SEQ ID NO:11492. Gaps Human genome, biallelic marker; high density disequilibrium map, genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer; ö / Match 0.8%; Score 13.8; DB 1; Length 21; Local Similarity 88.2%; Pred. No. 9e+02; les 15; Conservative 0; Mismatches 2; Indele 681 CACAGACAACCTTGTGG 697 BP. N 18 cacadacarccircies AAZ77136 standard; DNA; 21 10-SEP-2001 (first entry) diagnosis; ss AAZ77136; RESULT 1163 ò 셤

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Gaps

.; 0

2; Indels

cacadacarccrrcred 2

Homo sapiens W09954500-A2

28-OCT-1999

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99WO-IB000822

21-APR-1999;

Chumakov I;

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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haphotyphag studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3086, 3157, 3227, 3297 and 3037, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                          Novel biallelic markers used to construct a high density disequilibrium map of the human genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human biallelic marker downstream amplification primer SEQ ID NO:10380.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human genome, biallelic marker, high density disequilibrium map; genomic map, haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation, identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 13.8; DB 1; Length 21; Best Local Similarity 88.2%; Pred. No. 9e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 4 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                              Claim 9; Page 2680; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    364 GAGAGTGACCAGGCTTC 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAGAGTTACTAGGCTTC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ76024 standard; DNA; 21 BP
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98US-0109732P.
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                                98US-0082614P.
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                                                                                                                        Cohen D, Blumenfeld M,
                                                                                                                                                        WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention
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                                                                                   (GEST ) GENSET
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                                21-APR-1998;
23-NOV-1998;
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23-NOV-1998;
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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7440 represent amplification primers for the biallelic markers. The biallelic markers of the invention bave a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the pharmaceutical ascentical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3056, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, variant thrombospondin 1, variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forensics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                              Novel biallelic markers used to construct a high density disequilibrium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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/standard name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
iive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 7 A; 7 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human gene single nucleotide polymorphism #163.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bolk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ireland JS,
                                                                                               Claim 9; Page 2443; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1445 TGAAACATCCATTCTTC 1461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF95402 standard; DNA; 21 BP
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26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-025724P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 88.2
Matches 15; Conservative
                                                               of the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gargill M,
               WPI; 2000-013267/01.
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                                                                                                                                                                                                                                                                                                                                                                  present invention
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Variation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF95402;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various oblymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular the method can be used in the diagnosis of athersocherosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNDs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
     Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                         Query Match

0.8%; Score 13.8; DB 1; Length 21;
Best Local Similarity 88.2%; Pred. No. 9e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                           Sequence 21 3P; 6 A; 11 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human gene single nucleotide polymorphism #611.
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                                                                                                Example; Page 59; 242pp; English.
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26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-025724P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             850/c
AAF95850 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                     226 GAGAGTGGTGGTGGTGG
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                                                                   atherosclerosis.
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Variation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1166
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Mccarthy JJ;

Daley GQ,

Bolk S,

Gargill M, Ireland JS,

/*tag= a /standard_name= "single nucleotide polymorphism"

Location/Qualifiers replace(11,T)

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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various oblymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heat disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forensics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids comprising single nuclectide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
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                                                                                                                                                                                                                                                Sequence 21 BP; 5 A; 5 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human gene single nucleotide polymorphism #2182.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example, Page 198; 242pp; English
                              Example, Page 90; 242pp; English.
                                                                                                                                                                                                                                                                                                                                           190 AAGACCAATGGTGCCC 206
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВЪ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         replace (11, G)
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26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-025724P.
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                                                                                                                                                                                                                                                                                                                                                              AAF97421 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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atherosclerosis.
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Variation
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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
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                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                      Sequence 21 BP; 7 A; 8 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human gene single nucleotide polymorphism #1725.
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/*tag= a
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26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-025724P.
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                                                                                                                                                                                                                                                                                                                                392 CGGATGAGGTGCAGTCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                              15; Conservative
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Best Local Similarity
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Mccarthy JJ;

Bolk S, Daley GQ,

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in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPB) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, variant thrombospondin 1, variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
                                                                                                                                                                                                                      Gaps
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/standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                              0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
iive 0; Mismatches 2; Indels
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                                                                                                                                             Sequence 21 BP; 3 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human gene single nucleotide polymorphism #1343.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pulmonary embolism; paternity test; ds
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                                                                                                                                                                                                                                                       1031 CTGACTTTGGCCTGGCC 1047
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/*tag= a
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26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
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                                                                                                                                                                                                                  15; Conservative
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                                                                                                                                                                              Query Match
Best Local Similarity
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SNP specific lower PCR primer SEQ ID 3026.

(first entry)

14-AUG-2001

AAH40230;

AAH40230 standard; DNA; 21

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The present invention relates to a method for treating a patient diagnosed as having a lower than normal high density lipoprotein-cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, involving administering a compound that modulates LXR. or RXR-mediated transcriptional activity or ABC1 expression or activity. The LXR gene product may be used in an assay to identify compounds useful for the treatment of a disease or condition selected a lower than normal HDL cholesterol level, a higher than normal triglyceride level, and a cardiovascular disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating a lower than normal high density lipoprotein-cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, by administering a compound that modulates LXR- or RXR-mediated
useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                        High density lipoprotein-cholesterol; HDL-C; cardiovascular; ABCl; ds.
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                                                                                                   Score 13.8; DB 1; Length 21;
Pred. No. 9e+02;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 7 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
                                                                    Sequence 21 BP; 5 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 4; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                          Partial exon 7 corrected sequence.
                                                                                                                                                                             1268 CTGAGGAGACGTGGCCA 1284
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15-MAR-2000; 2000US-00526193.
23-JUN-2000; 2000US-0213958P.
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                                                                                                         0.8%;
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                                                                                                         Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
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Best Local Si
Matches 15;
                                                                                                                                                                                                                                                                                                                                        AAF93032;
                                                                                                                                                                                                                                                                  RESULT 1170
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Sequences AAH17205 - AAH40944 represent PCR primers, single nucleotide primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes Kits for determining the presence or absence of a SNP, using the oligomucleotides are useful for genotyping a genotyping primer. SNP flanking sequence, the SNP primer is used as a genotyping primer. SNP flanking sequence, the SNP primer is used as a genotyping primer. The oligomucleotides are useful for genotyping a mucleic acid sample by performing a single-nucleotide primer extension reaction. The cligomucleotides are useful for determining the presence, absence or identity of a SNP and for genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include disease e.g. caused by one or more SNPs. Phenotypic traits include disease e.g. caused by one or more SNPs. Phenotypic traits include disease e.g. caused by one or more SNPs. Phenotypic traits include disease.

CC caused by one or more SNPs. Phenotypic traits include disease e.g. craits also include symptoms of or suscoplania, polycystic kidney disease.

CC traits also include symptoms of or suscoplinity to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple solerosis, microorganism. The method is also useful in forensic investigations and microorganism. The method is also useful in forensic investigations and containing DNA sequence represents a PCR primer specific for a human SNP containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                             Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumacoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polymucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 6 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 65; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ORCH-) ORCHID BIOSCIENCES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0160096P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Picoult-Newburg L, Pohl M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-290930/30
                                                                                                                                                                                                                                                                                                                                                             WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-OCT-1999;
                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                        26-APR-2001.
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caddaagiricccidcr

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Gaps . 0

391

375 GGCTTCAGCCACGTCCT 17 decricascoascreer

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Conservative

Local Similarity es 15; Conserv

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Location/Qualifiers
                                                                                                                                                                                                                                                                                                Example 1; Page 20; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1480 ATCCACAAACTTCCTGA 1496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH89038 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          replace (11, t)
                                                                                    22-AUG-2000; 2000WO-US022986.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-NOV-2000; 2000WO-US030766.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0164596P,
                                                                                                                 23-AUG-1999; 99US-00379083.
07-JAN-2000; 2000US-00479128.
                                                                                                                                                              (DECO-) DECODE GENETICS EHF.
                                                                                                                                                                                          Gulcher J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17 AGCCTCAAACTTCCTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GLAX ) GLAXO GROUP LTD. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                       WPI; 2001-211306/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                          Olafsdottir BR,
                            WO200114555-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200134840-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-NOV-1999;
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-MAY-2001.
                                                         01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH89038;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4AH8903
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a purified and isolated non-naturally occurring DNA ligands to basic fibroblast growth factor (bFGF). The ligands are useful as part of gene therapy treatments and for diagnosing pathogenesis of vascular diseases including initiation and progression of atherosclerosis, acute coronary syndromes, vein graft disease and restencis following coronary angioplasty. The ligands have improved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, hypocretin receptor 1; orexin receptor 1; HCRTR1; chromosome 1; 1p33; central nervous system modulator; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel nucleic acid ligands to basic fibroblast growth factor that are useful as inhibitors of basic fibroblast growth factors and 2'-amino modified RNA ligands, exhibit increased in vivo stability.
                                                                                                                                                                         Ligand; basic fibroblast growth factor; bFGF; gene therapy; vascular; atherosclerosis; angioplasty; stability; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probe used to identify human hypocretin (orexin) receptor 1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ..
o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 13.8; DB 1; Length 21; Best Local Similarity 57.1%; Pred. No. 9e+02; Matches 12; Conservative 6; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 3 A; 4 C; 5 G; 2 T; 0 U; 7 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Col 69-75; 153pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CCGCGGCTCTGAGGTTGCTCG 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |:|:||| :|:| ||| CYGYGGCRYTRAARYTCCTCG 1
                                                       BP.
                                                                                                                                                                                                                                                                                                                                                       910S-00714131.
92US-00973333.
94US-00195005.
94US-00219012.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         160/c
AAF55160 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gold L, Tasset D;
                                                                                                                                                                                                                                                                                                               96US-00687421
                                                       AAF70928 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                   (NEXS-) NEXSTAR PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI, 2001-158583/16
                                                                                                                                               bFGF DNA ligand #61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         stability in vivo
                                                                                                                                                                                                                                                                                                                                                       10-JUN-1991;
06-NOV-1992;
10-FEB-1994;
                                                                                                                                                                                                                       Unidentified
                                                                                                                                                                                                                                                                                                               05-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-2001
                                                                                                                 20-APR-2001
                                                                                                                                                                                                                                                    JS6177557-B1
                                                                                                                                                                                                                                                                                                                                                                                                     28-MAR-1994;
                                                                                                                                                                                                                                                                                23-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Janjic N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF55160;
                                                                                   AAF70928;
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AAF55160, RESULT

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Probes AAFS5160-76 were used to identify a human hypocretin (orexin) receptor 1 (HCRTR1) gene. The HCRTR1 gene is present on chromosome 1, location 1p13. It is likely that a mutation in the HCRTR1 gene is associated with narcolepsy. HCRTR1 is a central nervous system modulator. The HCRTR1 polypeptide and polynucleotide are useful for diagnosing or treating narcolepsy in an individual. The HCRTR1 polynucleotide is a source of probes and primers, and is also used to produce the protein recombinantly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, single nucleotide polymorphic; SNP; forensic science; paternity testing; phenotypic trait; genetic mapping; animal breeding; plant breeding; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
Novel isolated nucleic acid molecule encoding hypocretin (orexin) receptor 1 useful for treating and diagnosing narcolepsy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
rative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human polymorphic oligonucleotide AC005336 fragment #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 6 A; 2 C; 8 G; 5 T; 0 U; 0 Other;
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The present invention relates to human oligonucleotides comprising a single nucleotide polymorphic site (SNP: AAH88797-AAH89219). The present sequence is one such oligonucleotide. The oligonucleotides can be used in forensics, paternity testing, correlation of polymorphisms with phenotypic traits, genetic mapping of phenotypic traits and marker assisted breeding of animals and crop plants
                                                                                    New polymorphic sites derived from the human genome are useful to determine sites correlating with phenotypic traits, particularly disease, and also in forensics and paternity testing.
                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 4 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
              Thomas D;
                                                                                                                                                                      Claim 71; Page 12; 43pp; English.
                ż
                Chen J, Patil
                Au K,
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Gaps
                                    .;
0
0.8%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 9e+02; rive 0; Mismatches 2; Indels
                    Local Similarity 88.2
nes 15; Conservative
        Query Match
                         Best Loca
Matches
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TGGTGCCCCTGAGCAGA 214 19 regaececreaecrea 3 198 Ġ 셤

RESULT 1175 ABA01349

ABA01349 standard; RNA; 21 ABA01349;

YMDD oligonucleotide #9. 03-JUL-2002

Selenoprotein; HIV; Ebola virus; cancer; immune system disorder; ss. Simian immunodeficiency virus.

JS6303295-B1 16-OCT-2001

Example 1; Page 44; 72pp; English.

95US-0001203P 14-JUL-1995; 01-SEP-1995;

2-JUL-1996;

(UYGE-) UNIV GEORGIA RES FOUND INC.

Ramanathan CS; raylor EW, Nadimpalli RG,

WPI; 2002-024734/03

New selenoprotein for use in detecting certain viruses, e.g. human immunodeficiency virus (HIV) or Ebola, cancer and immune system disorders

Disclosure; Col 69-70; 140pp; English

The present invention relates to selenoproteins encoded in the genome of a virus, where the coding sequence of the selenoprotein is genetically engineered for expression in a nucleic acid construct. The invention also discloses a method for identifying selenoprotein coding sequences, for detecting certain viruses (e.g. HIV or Ebola), cancer and immune system disorders. The present sequence was used to illustrate the invention

Sequence 21 BP; 7 A; 4 C; 4 G; 0 T; 6 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                      Probes for detecting target mucleotide sequence in sample, has sequence that forms hairpin structure having a double-stranded segment and singlestranded loop collectively forming region complementary to target
                Gaps
                ö
                Indels
Score 13,8; DB 1;
Pred. No. 9e+02;
3; Mismatches 2;
                                                                                                                                                          HPV; genotyping; nucleic acid detection; probe;
                                                                                                                                           DNA probe for human papilloma virus genotyping.
                                                                                                                                                                                                                                                                      (GENE-) APPLIED GENE TECHNOLOGIES INC.
                 3;
                                 866 AGCAGTACCTGGATGAC 882
                                                  21
                                                                                          ABA91520 standard; DNA; 21 BP
                                                                                                                                                                                                                              12-JUL-2001; 2001WO-US022166.
                                                                                                                                                                                                                                              14-JUL-2000; 2000US-00616761
30-MAR-2001; 2001US-00823647
0.8%;
                                            | |||:|| :|||:|||
5 ACCAGUACAUGGAUGAC
                                                                                                                            23-APR-2002 (first entry)
                 Conservative
                                                                                                                                                                            Human papillomavirus.
                                                                                                                                                                                                                                                                                                         WPI; 2002-171819/22.
         Local Similarity
ses 12; Conserv
                                                                                                                                                                                              WO200206531-A2.
                                                                                                                                                                                                                                                                                         Dattagupta N;
                                                                                                                                                                                                               24-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                  sequence.
  Query Match
            Best Loca
Matches
                                                                                             셤
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The present sequence comprises a probe for human papillomavirus (HPV)

C genotyping that was used in an example of the use of hairpin probes in

C mucleic acid hybridiaation analysis. The probe sequence is present within

the 5' stem portion of an RNA-DNA probe (see ABA91521) that is capable of

C forming a hairpin structure. The DNA protion of the hairpin probe

C includes methylphosphonates. The hairpin probe is immobilised onto a

membrane by BSA conjugation and the resulting probe-containing strip is

C contacted with HPV genomic DNA. After hybridised probe with RNA-DNA

structure. A second hybridisation is then performed using bictin-labelled

C probes, which are complementary to the portions of immobilised probe that

C become single-stranded after hybridisation and digestion. Biotin in the

C chemiluminescence. This is an example of the use of hairpin probes that

c are capable of both intrandolecular and internolecular hybridisation and

c n which the nucleotide sequence that is complementary to the target

C hairpin probe. The use of such probes reduces background hybridisation, ö Gaps .; 0 0.8%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 9e+02; iive 0; Mismatches 2; Indels Sequence 21 BP; 6 A; 8 C; 1 G; 6 T; 0 U; 0 Other; improving specificity 15; Conservative Local Similarity Query Match Matches

1677 CCCCAACTACATCTTCC 1693

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RESULT 1177

CCGTACTACATCTTCC 20

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Human; single nucleotide polymorphism; SNP; sickle cell anaemia; agammaglobulinaemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Miskott-Aldrich syndrome; Fabry's disease; familial hypercholesterolaemia; polycystic kidney disease, cancer; hereditary spherocycosis; Von Willebrand's disease; tuberous sclerosis; hereditary haemorrhagic telangiectasia; familial colonic polyposis; Ehlers-banlos syndrome; osteogenesis imperfects; autoimmune disease; acute intermittent porphyria; inflammation; nervous system disorder; infection; rheumatoid arthritis; multiple sclerosis; diabetes; systemic lupus erythematosus; draves disease; longevity; obesity; baldness; fertility; forensic; paternity testing; ss.
                                                                           Human single nucleotide polymorphism #97.
                                                                                                                                                                                                                                                                                                                                                                                                Cargill M, Ireland JS, Lander ES;
        ABK65477 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                        18-JAN-2001; 2001US-00765081.
                                                                                                                                                                                                                                                                                                                              19-JAN-2000; 2000US-0176861P.
                                                    02-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                  CARG/) CARGILL M.
IREL/) IRELAND J S.
LAND/) LANDER E S.
                                                                                                                                                                                                                                                                                                                                                                                                                     VPI; 2002-315108/35
                                                                                                                                                                                                                                                            US2002037508-A1.
                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                  8-MAR-2002
                               ABK65477;
ABK6547
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The invention relates to a nucleic acid comprising single nucleotide polymorphisms (SNPs) associated with diseases. The nucleic acids comprising the SNPs and probes and primers for detecting them may be used in assays for the diagnosis of diseases associated with SNPs (such as sickle cell anaemia, agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, mucular dystroppy, Wiskott-Addrich syndrome, Fabry's disease, familial hypercholesterolaemia, polycystic kidney disease, hereditary spherocytosis, Von Willebrand's disease, tuberous sclerosis, hereditary spherocytosis, Von Willebrand's disease, tuberous sclerosis, hereditary spherocytosis, Von Willebrand's disease, tuberous sclerosis, hereditary syndroms of or susceptibility to, multifactorial diseases of which a syndroms of, or susceptibility to, multifactorial diseases of which a component is or may be genetic, such as autoimmune diseases of which a inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms, autoimmune diseases including rheumatoid arthritis, multiple sclerosis, diabetes (insulin-dependent and non-carthritis, multiple sclerosis, diabetes (insulin-dependent), strength, speed, endurance, fertility, ovary, pancreas, prostate, skin, scomach and uterus, longevity, appearance (e.g., baldness, cobesity), strength, speed, endurance, fertility, and susceptibility or receptivity to particular drugs or therapeurometer treatments), in forensics and in paternity testing. Nucleic acid comprising single nucleotide polymorphisms, useful in forensics, paternity testing and diagnosis of disease. Claim 1; Page 46; 96pp; English.

Sequence 21 BP; 4 A; 4 C; 4 G; 8 T; 0 U; 1 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; Clesterase inhibitor; ClNH; kallikrein 1; XRX1; bradykinin receptor B2; BDKRB2; gene therapy; and the receptor B2; BDKRB2; gene therapy; and converting enzyme 2; ACB2; protease inhibitor 4; P14; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; thugal infection; COPD; Chronic obstructive pulmonary disease; enterocolitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an isolated nucleic acid from a human gene encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (BDXBB1), tachykinin receptor B1 (TACR1), C1 esterase inhibitor (C1NH), kallikrein (KAK1), bradykinin receptor B2 (BDXRB2), angiotensin converting enzyme 2 (ACE2) or protease inhibitor 4 (P14), comprising at least one polymorphic position. Also included are (1) a probe that hybridises to a polymorphic position as provided in the detailed summary of single nucleotide polymorphisms comprising additional 5' and 3' flanking genomic sequence; (2) analysing (M1) at least one nucleic acid sample comprising obtaining the sample from one or more individuals and determining the mucleic acid sequence at one or more polymorphic positions in a gene encoding a protein selected from the group above; (3) constructing (M2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful
                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated nucleic acid with at least one polymorphic position, use for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Perrone MH;
                                                                  Indels
DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ma-Edmonds M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human polymorphism associated DNA sequence #445.
Score 13.8; DB 1
Pred. No. 9e+02;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 883; 977pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zerba KE,
                                                                                                                                     767 TCAAGGACCTCAAACACGC 785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BRIM ) BRISTOL-MYERS SQUIBB CO. (TSUC/) TSUCHIHASHI Z. (HUIL/) HUI L.
                                                                                                                                                                             TCAAAGATGTYAAACACGC 3
                                                                                                                                                                                                                                                                                                                                                                             ABS60808 standard; DNA; 21 BP.
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23-JAN-2001; 2001US-0263678P.
02-MAR-2001; 2001US-0273037P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-DEC-2001; 2001WO-US047235.
Query Match
Best Local Similarity 78.9%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-NOV-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ŗ,
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Swanson BN, Powell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-619265/66.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABS60808;
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haplotypes using the genes comprising grouping at least two nucleic acids (4) identifying (M3) an individual at risk of developing a disorder upon administration of an ACE inhibitor and/or vasopeptidase inhibitor using the polymorphic data; (5) a library of nucleic acids, each of which comprises one or more polymorphic positions within a gene encoding a human protein selected from the group above; and (6) genotyping (M4) an individual comprising obtaining a mucleic acid sample, determining the nucleotide present in at least one polymorphic position, and comparing at least one position with a known data set. The genes, (M1, M2, M3 and M4) and compositions are useful for detecting, disgnosing, treating, preventing various disorders such as angloedaema and diseases which involve anglogenesis like haemangiomas, tumours, sarcomas, Crohn's involve anglogenesis like haemangiomas, tumours, sarcomas, Crohn's involve anglogenesis like haemangiomas, tumours, sarcomas, Crohn's hypertension, heart failure, myocardial infarction, ventricular hypertension, heart failure, myocardial infarction, ventricular hypertension, heart failure, myocardial infarction, ventricular hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory artery disease, and and/or atherosclerosis and/or artery disease, and disorders are listed in the specification). The obstructive pulmonary disease (COPD) and enterocollitis (many other diseases and disorders are listed in the specification) and bloodies are also useful for chromosome identification. Antibodies and biological samples. The present sequence is included in the sequence and biological samples. The present sequence is included in the sequence in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein 1; KLR1; bradykinin receptor B2; BDKRB1; gene therapy; and addykinin receptor B2; BDKRB1; gene therapy; and converting enzyme 2; ACR2; protease inhibitor 4; P14; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; coronary artery disease; angloedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD; chronic obstructive pulmonary disease; enterocolitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.8; DB 1; Length 21;
Pred. No. 9e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 6 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 fuman polymorphism associated DNA sequence #332.
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(TSUC/) TSUCHIHASHI Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                915 ACTGTTCCTGTTCCAGC 931
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23-JAN-2001; 2001US-0263678P.
02-MAR-2001; 2001US-0273037P.
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Best Local Similarity 88.2.
Thes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1179
ABS60583/c
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0; Gaps

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The invention relates to an isolated nucleic acid from a human gene encoding aminopeptidase P (KRNEP2), bradykinin receptor B1 (TACR1), C1 esterase inhibitor (CINH), kallikrein tachykinin receptor B1 (TACR1), C1 esterase inhibitor on the care of the control of the care 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polymucleotides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence
                                                                                                                                                     New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    listing but is not referred to anywhere else in the specification
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                                          Perrone MH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
ative 0; Mismatches 2; Indels
                                       Ma-Edmonds M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 6 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human polymorphism associated DNA sequence #331.
                                                                                                                                                                                                                                                                                Disclosure, Page 809, 977pp; English.
                                          Zerba KE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             915 ACTGTTCCTGTTCCAGC 931
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                                          Swanson BN, Powell JR;
                                                                                                           WPI; 2002-619265/66.
                                                                                                                                                                                                                                autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 15; Conserv
(HUIL/) HUI L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15;
                                                                     Swanson BN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABS60582;
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ABS60582/c
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04-DEC-2000; 2000US-0251015P. autoimmune diseases. WO200261131-A2. (TSUC/) TSUCHI (HUIL/) HUI L. Fsuchihashi Z, Homo sapiens. 08-AUG-2002 Swanson BN,

tachykinin receptor B1, TACR1, C1 esterase inhibitor; C1NH; kallikrein 1, KLK1; bradykinin receptor B2; BDKB2; gene therapy; angiotensin converting enzyme 2; ACE2; protease inhibitor 4; P14; polymorphism, heamangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardal infarction; ventricular hypertrophy; vascular disease; angioedaema; arteriosclerosis; anherosclarosis; connary artery disease; angioedaema; arteriosclerosis; atherosclarosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD; Chronic obstructive pulmonary disease; enterocolitis.

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03-DEC-2001; 2001WO-US047235.

23-JAN-2001; 2001US-0263678P. 02-MAR-2001; 2001US-0273037P.

(BRIM) BRISTOL-MYERS SQUIBB CO. (TSUC/) ISUCHIHASHI Z.

Zerba KE, Ma-Edmonds M, Perrone MH; Z, Hui L, Powell JR;

WPI; 2002-619265/66.

New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and

Disclosure; Page 809; 977pp; English.

The invention relates to an isolated nucleic acid from a human gene encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (BDXRB1), cleatezase inhibitor (CMR1), kallikrain teceptor B1 (TACR1), cleatezase inhibitor (CMR1), bradykinin receptor B2 (BDXRB2), angiotensin converting enzyme (CMLKA1), bradykinin receptor B2 (BDXRB2), angiotensin converting enzyme 2 (ACE2) or protease inhibitor 4 (P14), comprising at least one paymorphic position. Also included are (1) a probe that hybridises to a polymorphic position as provided in the detailed summary of single curl coloring polymorphisms comprising additionals and 3' flanking genomic sequence; (2) analysing (M1) at least one nucleic acid sequence at one or more polymorphic positions in a gene cuccing a protein selected from the group above; (3) constructing (M2) and polymorphic positions within a specencial adjocation and individual at risk of developing a disorder using the polymorphic data; (5) albrary of nucleic acids, each of which comparises one or more polymorphic positions within a spece encoding a comprises one or more polymorphic positions within a specencial of the polymorphic positions within a specencial of protein selected from the group above; and (6) genotyping (M4) and individual comparising at least one polymorphic position, and comparing at least one polymorphic position, and comparing the present in at least one polymorphic position, and comparing the present in at least one polymorphic position, thrombosis, coronary and compositions are useful for detecting, diagnosing, treating, and sease, trachomas, and cardiovascular diseases like angine pectoris, hypertrophy, vascular diseases, aneurysm, embolism, thrombosis, coronary artery disease, arterioral arterioral and engale and diseases which are polymorphic position, heart failure, myocardial infarction, ventricular hypersensitivity reactions, sepals, autoimmune diseases, inflammary disease, are listed in the specification). The polymucleotides are also useful for chromosome identification).

The invention relates to a novel human antibody (I), preferably a human monoclonal antibody which binds to an activation inducible lymphocyte immunoadulatory molecule (ALILM). (I) is useful for modulating signal transduction into a cell mediated by ALILM, for modulating proliferation of AILM-expressing cells, for modulating production of a cytokine from AILM-expressing cells, and for inducing antibody-dependent cytotoxicity against AILIM-expressing cells and/or immune cytolysis or apoptosis of AILM-expressing cells and/or immune cytolysis or apoptosis of AILM-expressing cells and/or immune cytolysis or apoptosis of Prophylaxis of delayed type allergy. (I) is useful for treating, preventing or preventing various diseases associated with AILIM-mediated costimulatory transduction, and for inhibiting the onset and/or advancement of the diseases. (I) is useful for suppression, prevention and/or treatment of

New human monoclonal antibody that binds to activation inducible lymphocyte immunomodulatory molecule, useful for treating rheumatoid arthritis, multiple sclerosis and inflammation.

Example 10; Page 247; 300pp; English.

ö against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence listing but is not referred to anywhere else in the specification Human, antirheumatic; antiarthritic; antidiabetic; antipsoriatic; antiallergic; antiulcer; neuroprotective; antithyroid; vasotropic; immunosuppressive; dermatological, antiinfiammatory; hepatorropic; activation inducible lymphocyte immunomodulatory molecule; AILIM; monoclonal antibody; allergy; rheumatoid arthritis; diabetes mellitus; multiple selerosis; autoimmune thyroiditis; psoriasis; hepatitis; allergic contact-type dermatitis; chronic inflammatory dermatosis; systemic lupus erythematosus; autoimmune disorder; inflammation; graft versus host reaction; immune disorder; intestinal immunity; pCR primer; ss. Gabs ö Query Match

0.8%; Score 13.8; DB 1; Length 21;
Best Local Similarity 88.2%; Pred. No. 9e+02;
Matches 15; Conservative 0; Mismatches 2; Indels Anti-human AILIM monoclonal antibody, sequencing primer #2. Sequence 21 BP; 6 A; 5 C; 6 G; 4 T; 0 U; 0 Other; 915 ACTGTTCCTGTTCCAGC 931 AAS99452 standard; DNA; 21 BP. 19 Acrerreredec 3 15-MAY-2001; 2001WO-JP004035. 18-MAY-2000; 2000JP-00147116. 30-MAR-2001; 2001JP-00099508. Tsuji T, Tezuka K, Hori N; 12-MAR-2002 (first entry) (NISB) JAPAN TOBACCO INC. WPI; 2002-075313/10. WO200187981-A2. Homo sapiens. Synthetic. 22-NOV-2001. AAS99452; RESULT 1181 AAS99452 ន្តន្តន្តន

sequence is used to illustrate the method of the invention

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rheumatoid arthritis, multiple sclerosis, autoimmune thyroiditis, allergic contact-type dermatisis, chronic inflammatory dermatosis, systemic luques erythematosus, insulin-dependent diabetes mellitus, psoriasis, autoimmune or allergic disorders, inflammation, graft versus host reaction, graft versus host reaction, graft versus host disorders immune rejection, disorders intestinal intestinal inmunity, specifically inflammatory intestinal disorders such as ulcerative colitis, pneumonia, hepatitis, nephritis, vasculitis, and pancreatitis. (I) induces no serious antimunorejection due to antigenicity to human, i.e., human anti-mouse antigenicity (HAMA) in a host. AASS9444+AASS9477 represent anti-human ALLIM monoclonal antibody coding sequences and PCR primers of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunogenic composition of mycobacterial mutants with modified protein production capabilities, useful for vaccinating and treating infections in particular mycobacterial diseases such as tuberculosis and Crohn's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunogenic; infection; vaccine; mycobacterial disease; tuberculosis;
Crohn's disease; gene therapy; antiinflammatory; antibacterial; hsp70;
heat shock protein 70; PCR; primer; ss.
                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mycobacterium sp. hsp70 operon promoter amplifying primer, Hsp701.
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                                                                                                                                                                                                                                              Score 13.8; DB 1; Length 21;
Pred. No. 9e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                Sequence 21 BP; 4 A; 4 C; 9 G; 4 T; 0 U; 0 Other;
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Best Local Similarity 88.2%;
Matches 15; Conservative
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29-MAY-2001; 2001US-0294170P.
                                                                                                                                                                                                                                                                                                             849 CCTGGACAAGGACCTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SEQU-) SEQUELLA INC.
(YOUN/) YOUNG D B.
(STEW/) STEWART G R.
(OGAO/) O'GAORA P C E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stewart GR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-698637/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mycobacterium sp.
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                                                                                                                                                                                  invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD45724;
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The invention relates to an immunogenic composition of mycobacterial mutants with modified protein production capabilities. The invention also relates to methods for the treatment and prevention of infections also diseases. The methods and compositions of the invention are useful for vaccinating and treating infections in particular mycobacterial diseases such as tuberculosis and Crohn's disease. The invention is also used in gene therapy. The present sequence is a PCR primer used for amplifying Mycobacterium sp. hsp70 (heat shock protein 70) operon promoter. This

Example 9; Page 29; 59pp; English.

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The present invention relates to a method of diagnosing a cellular proliferative disorder of breast tissue, which involves determining the state of methylation of one or more nucleic acids isolated from the state of methylation of the nucleic acids acompared with a state of methylation from a subject not having the cellular proliferative disorder of breast tissue is indicative of a cellular proliferative disorder of breast tissue is indicative of a cellular may be TWIST, HOXAS, NES-1, retinoic acid receptor beta (RARbeta), oestrogen receptor, cyclin D2, Wilms' tumour gene (WT-1), 14.3.3 sigma, HN-10 r RASSFR. The method is useful for diagnoshig and/or determining a predisposition to a cellular proliferative disorder, in particular breast cancer including ductal carcinoma in situ, lobular carcinoma, colloid carcinoma, tubular carcinoma, medullary carcinoma, metaplastic carcinoma, intraductal carcinoma, medullary carcinoma in situ and papillary carcinoma in situ. The present sequence is a primer used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proliferative disorder of breast tissue, in particular breast cancer, by determining the state of methylation of one or more nucleic acids isolated from the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                 Human; methylated gene; methylation; breast cancer; marker; WT-1; cell proliferative disorder; TWIST; HOXAS; NES-1; RARbeta; cyclin D2; retinoic acid receptor beta; oestrogen receptor; Wilms' tumour; 14.3.3 sigma; HIN-1; RASSFIA; tumour suppressor gene; hypermethylation;
                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Davidson N, Fackler MJ;
                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diagnosing and/or determining a predisposition to a cellular
                                                                      Score 13.8; DB 1; Length 21;
Pred. No. 9e+02;
); Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.8; DB 1; Length 21; Pred. No. 9e+02;
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                                     Sequence 21 BP; 4 A; 5 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dooley WC, Sacchi N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE.
                                                                                                                                                                                                                                                                                                                                                                                    Cyclin 14-3-3 sigma gene PCR primer #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 12; Page 44; 115pp; English.
                                                                                                         .
                                                                                                                                              1020 GCTCAAGCTGGCTGACT 1036
                                                                                                                                                                                 3 GGTCAAGCTGGCGGACT 19
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                                                                        0.8%;
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                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                            Query Match
Best Local Similarity 88.2'
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Best Local Similarity
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                    843 TGAGTACCTGGACAAGG 859
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15; Conservative
Matches
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RESULT 1184

ABS97470 standard; DNA; 21 ABS97470

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ABS97470;

(first entry)

23-DEC-2002

Human diazepam binding inhibitor (DBI) gene polymorphic sequence #14.

Human, ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;

Cytochrome P450 A2; CYP4501A1; CYP601B2; CYP4501B1; LTF;

Cytochrome P450 A2; CYP4501A2; CYP601CEE CYP45002E1; LTF;

Cytochrome P450 A2; CYP4501A2; CYTOCHROME P450 02E;

Cytochrome P450 A2; CYP4501A2; CYTOCHROME P481; MRP3; NR112;

Cytochrome Parceptor nuclear translocator; ARNT, cathepsin S; CTSS;

Cytochrome Parceptor nuclear translocator; ARNT, cathepsin S; CTSS;

Cytochrome Parceptor nuclear translocator; ARNT, cathepsin S; CTSS;

Cytochrome Parceptor nuclear transferase;

CYCLOX2; diazepam binding inhibitor; DB1; haematological;

CYCLOXCHROME PARCEPTOR PARCEPTOR PARCEPTOR PLAP;

CYCLOXCHROME PARCEPTOR PARCEPTOR PARCEPTOR PLAP;

CYCLOXCHROME PARCEPTOR PARCEPTOR PARCEPTOR PLAP;

CYCLOXCHROME CANAGE PARCEPTOR PARCEPTO single nucleotide polymorphism.

Homo sapiens.

25-JUL-2002.

28-NOV-2001; 2001WO-US044838.

28-NOV-2000; 2000US-00724389.

DNAS-) DNA SCI LAB INC.

Guida M, Hall J;

WPI; 2002-698522/75.

Isolated nucleic acid molecules having polymorphisms in known human genes eg. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits.

Example 9; Page 115; 714pp; English.

This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known molecule comprising at least one base variation from that of a known wolconome P450 A1 (CY45510A1), atcremented A2 (CY4551A2), cytochrome P450 D2E1 (CY4550D2E1), adrenergic receptor beta1 (ADDR1), aryl hydrocarbon receptor nuclear translocator (ARNT), cathepsin 8 (CYS2), cytloxygenase 2 (GXOZ), diazepam binding inhibitor (DBI), epoxide hydroxylase 2 (EMXZ), 5-lipoxygenase activating protein (FLAP), glutathione-S-transferase 12 (GXI12), histamine-N-methyl transferase (HNWT), NADPH quinone oxidoreductase 2 (NQOZ), sulfotransferase (HNWT), NADPH quinone oxidoreductase 2 (NQOZ), sulfotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl transferase (UGT2B1), urokinase receptor (URPS), multidrug resistance 1 (MDR1), actotransferzin (LTF), unltidrug resistance associated protein 3 (MRP3), orphan nuclear receptor (NRIIZ), or acetylcholine muscarinic

The polymorphisms in the human genes cited in the invention are useful as genetic linkage markers for locating and characterising the genes that are responsible for specific traits within the genome and eventually identifying the genes responsible for a variety of disorder-related trains as a result of thair e.g., overexpression, constitutive expression, mutation or underexpression, which may be used in diamosing and/or treating the disorders. The nucleic acid molecules comprising the polymorphic sequences contained in CYP4501A1, CYP4510A2, ARNY, BEHX, GST12, NNMY, NQC2, NR112, STW, UGT2B4, UGT2B5, AHR, MDR1 and/or MDR3 are useful for screening individuals for altered drug metabolism. The polymorphic sequences contained in CYP4501A1, CYP4501A2, ARNY, MDR1 and/or MDR3 may also be used to screen individuals for altered central used to screen for altered cardiovascular function, in COX2 for altered central nervous system function, in FADP and HNWY for altered pulmonary, immological the metabological function, in FADP and HNWY for altered serine function, in FADP and HNWY for altered immunological or haematological function, in FADP and HNWY for altered central and peripheral nervous system function, in CHNR3, CHNR4 for Altered central and peripheral nervous system function. The present sequence represents a polymorphic DNA sequence of the invention

Sequence 21 BP; 6 A; 6 C; 6 G; 3 T; 0 U; 0 Other;

Gaps ö Query Match 0.8%; Score 13.8; DB 1; Length 21; Best Local Similarity. 88.2%; Pred. No. 9e+02; Matches 15; Conservative 0; Mismatches 2; Indels

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704 AGGAGATCAGACTGGAA 720 21 s accadercadacredaa

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1185 ABK53783

BP. ABK53783 standard; DNA; 21

ABK53783;

(first entry) 05-JUN-2002

DMS:acceptor oxidoreductase, PCR primer #29.

DMS:acceptor oxidoreductase, dimethyl sulphide, sulphoxide, prochiral organic sulphide; sulphoxide enantiomer; primer; chiral drug production; optically-active functional drug; ss.

Rhodovulum sulfidophilum

WO200216570-A1.

28-FEB-2002.

21-AUG-2001; 2001WO-AU001033.

21-AUG-2000; 2000AU-00009559.

(UYQU) UNIV QUEENSLAND.

Mcewan AG; Mcdevitt CA,

WPI; 2002-280922/32.

New recombinant dimethyl sulfide:acceptor oxidoreductase or its subunits, useful for oxidizing prochiral organic sulfides to form sulfoxide enantiomers for chiral drug synthesis.

Claim 15; Page 46; 66pp; English.

The invention relates to a recombinant dimethyl sulphide (DMS):acceptor oxidoreductase (I) or its subunit selected from recombinant alpha, beta, delta and gamma subunits. (I) is useful for oxidising prochiral organic

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The invention describes a polymuclectide (I) of the endothelin (EDN)/endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) signaling system which is associated with a cardiovascular disease. (I), the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) or (III) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (II), (III), the EDN, ECE or EDNR FOLYPEPTIGE, or a cardiovascular disease. (II), till), the EDN, ECE or EDNP polypeptide, or a cardiovascular disease obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE
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               is expressed in a transformed bacterium. The enantiomer formed is useful actor producing a chiral drug. (I) is useful for spribesis of optically-active functional groups of drug. DNA encoding (I) is useful for producing a strain of DMS:acceptor oxidoreductase- deficient Rhodovulum sulfidophilum, which is useful in whole-cell reaction, where DMS:acceptor oxidoreductase activity is unwanted. ABK53751-ABK53805 represent R. sulfidophilum DMS:acceptor oxidoreductase subunit coding sequences and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angliogenesis; farty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.
to form sulphoxide enantiomers for chiral drug synthesis. (I)
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                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 9e+02; ive 0; Mismatches 2; Indels
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signaling system or its gene product. The isolated proteins and polynucleotides encoding them are useful for preparation of a pharmacutical composition for treating a cardiovascular disease such as coronary heart disease, hypertension, atherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercholesterolaemia. The gene or a polynucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support comprising polynucleotides, genes, vectors, polypeptides, antibodies or host cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angiogenesis; tatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Endothelin converting enzyme 1 (ECE-1) SNP detection primer #143.
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                                                                                                                                                                                                                                                                                                                0.8%; Score 13.8; DB 1; Length 21;
88.2%; Pred. No. 9e+02;
iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                           Sequence 21 BP; 5 A; 7 C; 9 G; 0 T; 0 U; 0 Other;
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Matches 15; Conservative
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molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and polymucleotides encoding them are useful for preparation of a polymucleotides encoding them are useful for preparation of a pharmaceutical composition for treating a cardiovascular disease such as coronary heart disease, hypertension, atherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercholesterolaemia. The gene or a polymucleotide fragment of the CEN/EE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support comprising polymucleotides, genes, vectors, polypeptides, antibodies or host cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; interleukin 4 receptor; IL4R; type 1; diabetes; allele; insulian dependent diabetes mellitus; IDDM; myasthenia gravis; single nucleotide polymorphism; SNP; autoimmune disease; T helper type 1 mediated disease; rheumatoid arthritis; probe; multiple solerosis; inflammatory bowel disease; systemic solerosis; systemic lupus erythematosus; psoriasis; soleroderma; Grave's disease; Guillain-Barre syndrome; Hashimoto's thyroiditis; so.
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88.2%; Pred. No. 9e+02;
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mellitus (IDDM)-associated interleukin 4 receptor allele in a nucleic acid sample of the individual, where the presence of the allele indicates the individual's risk for type I diabetes. The method identifies one or more single nucleotide polymorphism (SNP) within the ILMR gene listed in the specification. The method and the SNP's are useful for determining an individual's risk for type I diabetes. The ILMR SNPs are also useful for determining an individuals risk for any autoimmune disease or condition or any T helper type I mediated disease, e.g. rheumatoid arthritis, multiple sclerosis, inflammatory bowel disease, systemic lupus sclerosis, mysathenia gravis, Guillain-Barre syndrome, or Hashimoto's thyroiditis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Determining an individual's risk for type 1 diabetes, comprises detecting the presence of an insulin dependent diabetes mellitus-associated interleukin 4 receptor allele in a nucleic acid sample of the individual.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequences given in ABQ80153-69 represent probes which were used to identify wild type and variant loci in the human interleukin 4 receptor (ILAR). These probe sequences were used in the method of the invention for determining an individual's risk for type 1 diabetes. The method comprises detecting the presence of an insulin dependent diabetes andlitus (IDDM) associated interleukin 4 receptor allele in a mucleic acid sample of the individual, where the presence of the allele indicates the individual's risk for type 1 diabetes. The method identifies one or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; interleukin 4 receptor; ILAR; type 1; diabetes; allele; insulin dependent diabetes mellitus; IDDM, wyasthenia gravis; single nucleotide polymorphism; SNP; autoimmune disease; T helper type 1 mediated disease; rheumatoid arthritis; probe; multiple sclerosis; inflammatory bowel disease; systemic sclerosis; systemic lupus erythematosus; psoriasis; scleroderma; Grave's disease; guillain-barre syndrome; Hashimoto's thyroiditis; ss.
                                                                                                                                                                                                                                                                                                                                                       / Match 0.8%; Score 13.8; DB 1; Length 21; Local Similarity 88.2%; Pred. No. 9e+02; les 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mirel DB, Erlich HA, Bugawan TL, Noble JA, Valdez AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Probe DBM0080P, identifies wild type IL4R SNP #8.
                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 3 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HOFF ) ROCHE DIAGNOSTICS GMBH. (HOFF ) HOFFMANN LA ROCHE & CO AG F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; Page 36; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         1175 TCTTCTATGAGATGGCC 1191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 rcrrcrcraacaraccc 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-JUL-2002; 2002WO-EP007956.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-JUL-2001; 2001US-0306912P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABQ80161 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUN-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABQ80161;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1189
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     8888888888888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
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G; 6 T; 0 U; 0 Other;

Sequence 21 BP; 6 A; 8 C; 1

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Query Match
888888888888
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more single nucleotide polymorphism (SNP) within the IL4R gene listed in the specification. The method and the SNP's are useful for determining an individual's risk for type 1 diabetes. The IL4R SNP's are also useful for determining an individual's risk for any autoimmune disease or condition or any Thelper type 1 mediated disease, e.g. rheumatoid arthritis, multiple sclerosis, inflammatory bowel disease, systemic lupus schematosus, psoriasis, scleroderma, Grave's disease, systemic thynsolatis, myasthenia gravis, Guillain-Barre syndrome, or Hashimoto's thyroiditis
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Sequence 21 BP; 3 A; 7 C; 5 G; 6 T; 0 U; 0 Other;

DB 1; Length 21; 2; Indels Score 13.8; DB 1 Pred. No. 9e+02; 0; Mismatches ; 0 1175 TCTTCTATGAGATGGCC 1191 0.8%; ilarity 88.2%; Conservative Local Similarity es 15; Conserv Matches ò

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Gaps

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rcrrcrcradaraccc 18 g

AAL53951 standard; DNA; 21 AAL53951; RESULT 1190

BP

(first entry) 18-FEB-2003

Human papillomavirus probe, SEQ ID No 1.

hybridising; target DNA; duplex; RNase H; Detecting; point mutation, hybridising; ta single nucleotide polymorphism; probe; ss.

Human papillomavirus

US2002142308-A1

30-MAR-2001; 2001US-00823634

10-MAR-2001; 2001US-00823634

(DATT/) DATTAGUPTA N. (TSEN/) TSENG T.

WPI; 2003-102506/09

Dattagupta N, Tseng T;

Detecting point mutation in DNA strand, by hybridizing target DNA strand having mutation with test DNA strand to form duplex, contacting the duplex with RNase H and determining the cleavage of test strand by RNase

Example 1; Page 12; 26pp; English.

The invention relates to a novel method for detecting a point mutation in a DNA strand. The novel method comprises hybridishing a target DNA strand containing or suspected of containing a point mutation with a test nucleic acid strand complementary to the DNA strand to form a target DNA strand/test nucleic acid strand duplex, contacting the duplex with an strand/test nucleic acid strand duplex, contacting the duplex with an enclosive H, and determining whether the ribonucleotide residues within the nucleotide sequence are cleaved by RNase H. The method is useful for detecting a point mutation in a DNA strand, where the point mutation to be detected is a single nucleotide polymorphism, preferably a colymorphism in a genome, e.g., a viral, bacterial, eukaryotic, mammalian or human genome. The method is useful to detect any nucleic acids from any species of organisms such as Acintobacter, Bacillus, Candida, Enterococcus, Heemophilus, Mycobacterium and Streptococcus, and viruses. This polymucleotide sequence represents a probe relating to the mutation detecting method of the invention

Human stearyl coenzyme A desaturase 4 siRNA Desat3-antisense.

ADC72204 standard; RNA; 21

1192

RESULT 1. ADC72204,

ADC72204;

BXZXXXB

18-DEC-2003 (first entry)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Activation of potential matrix metalloprotease-2 (proMMP-2) with claudins via membrane type matrix metalloproteases (MT-MMPs).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention describes the activation of potential matrix metalloprotease-2 (proMMP-2) with claudins via membrane type matrix metalloproteases (MT-MMP). Activated proMMP-2 is useful for treatment neovascularisation and cancer. This sequence represents a potential matrix metalloprotease-2 activation associated primer. Note: This sequence is given in the specification as seq id 13.
                                                                                                                                                                                                                                                                                vasotropic; cytostatic; potential matrix metalloprotease-2; proMMP-2; membrane type matrix metalloproteases; MT-MMP; neovascularisation; cancer; human; claudin 1; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                  Gaps
                                                                                                                                                                                                                                                         Potential matrix metalloprotease-2 activation related primer seq id
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88.2%; Pred. No. 9e+02;
ve 0; Mismatches 2; Indels
Score 13.8; DB 1; Length 21;
Pred. No. 9e+02;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; SEQ ID NO 31; 49pp; Japanese.
                                                                1693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          453 CACTGAGGACATCAACA 469
                                                                                           4 CCGTAACTACATCTTCC 20
                                                                                                                                                       528/c
ADC51528 standard; DNA; 21 BP.
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 0.8%;
Local Similarity 88.2%;
Nes 15; Conservative
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                                                               1677 CCCCAACTACATCTTCC
                                                                                                                                                                                                                           (first entry)
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les 15; Conserv
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                                                                                                                                                                                                                                                                                                                                               Synthetic.
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   Query Match
                                                                                                                                                                                              ADC51528
                                                                                                                                       RESULT 1191
                                 Matches
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Matches
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This invention relates to a novel method of identifying candidate p53
pathway modulating agent which comprises contacting an assay system
comparising stearyl coenzyme A desaturase (SCD) polypeptide or nucleic
agent under conditions where, but for the presence of the test agent, the
system provides a reference activity. The p53 gene is mutated in over 50
different types of human cancer and is believed to be the most commonly
mutated gene in human cancer. The human p53 protein normally functions as
a central integrator of signals including DNA damage, hypoxia, nucleotide
optivation and oncogene activation. Modulators of p53 may have
carrial integrator of signals including DNA damage, hypoxia, nucleotide
optivation and oncogene activation. Modulators of p53 may have
invention are useful for identifying modulators of p53 may have
therapeutic targets for disorders associated with defective p53 function,
such as anglogenic, apoptotic or cell proliferative disorders, for
cample cancer. The modulators are useful as research reagents,
clasmostics and therapeutics. The present sequence (see ) small
chat of a human Stearoyl-CoA desaturase 4 sequence (see ) small
interfering RNA (siRNA) used in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying a candidate p53 pathway-modulating agent as therapeutic targets for disorders related to defective p53 function e.g. cancer by contacting an assay system having SCD polypeptide or nucleic acid, with a
                  p53 pathway modulating agent; stearyl coenzyme A desaturase; SCD; p53 gene; cancer; signal integrator; DNA damage; hypoxia; nucleotide deprivation; oncogene activation; cytostatic; gene therapy; defective p53 function; angiogenic; apoptotic; cell proliferative disorder; human; small interfering RNA; siRNA; ss; Desat3-antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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primer, haplotype, familial glucocorticoid deficiency; FGD, cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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Pred. No. 9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                    Friedman L, Plowman GD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 5 A; 6 C; 4 G; 0 T; 4 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; SEQ ID NO 30; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1637 GGCAGCGGCTGGAGGGATGCC 1657
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                                                                                                                                                                                                                                                                28-FEB-2003; 2003WO-US006087.
                                                                                                                                                                                                                                                                                                         01-MAR-2002; 2002US-0361196P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%;
                                                                                                                                                                                                                                                                                                                                                                                  Francis-Lang H,
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                                                                                                                                                                                                                                                                                                                                           EXEL-) EXELIXIS INC
                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-748276/70.
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Best Local Similarity
                                                                                                                                                                                         WO2003074662-A2.
                                                                                                                                                    Homo sapiens.
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                                                                                                                                                                                                                             12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                    Belvin M,
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The present invention provides the gene, protein and cDNA sequences of the human melanocortin 2 receptor (adrenocorticotrophic hormone) (WC2R). Also identified are a number of single nucleotide polymorphisms (SNPS) found within the sequences. The sequences can be used to find the haplotype of the MC2R gene in an individual and to identify drugs for the reatment of cancer and familial glucocorticoid deficiency. The present sequence is an allele specific primer for the gene of the invention, which is found on chromosome 18q11.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 constituent of gag gene of HIV-1 Bru, -Mal or -Eli, HIV-
                                                                                                                                                                                                                                                  Melanocortin 2 receptor (MC2R) gene polymorphic variants, useful e.g. in studying the expression and function of MC2R and screening candidate drugs for treating familial glucocorticoid deficiency and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.6; DB 1; Length 15;
Pred. No. 7e+02;
1; Mismatches 0; Indels
chromosome 18q11.2; SNP; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV-1; HIV-2; SIV; AIDS; anti-sense nucleotide; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 2 A; 4 C; 5 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                               Lee HH, Sausker EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (INSP ) INST PASTEUR.
(INRM ) INSERM INST NAT SANTE RE.
                                                                                                                                                                                                                                                                                                           Claim 16; Page 14; 79pp; English.
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                                                                                                                                                                    (GENA-) GENAISSANCE PHARM INC.
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                                                                                                                                        30-JUN-2000; 2000US-0215330P.
                                                                                                             29-JUN-2001; 2001WO-US021064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8%;
Best Local Similarity 92.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGTTCTGCACGGGK 14
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13; Conservative
                                                                                                                                                                                                                         WPI; 2002-171650/22.
                                                                                                                                                                                              Koshy B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MMy4B nucleotide of ROD and SIV-MAC.
                                                       WO200202821-A1
                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUN-1990;
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                                                                                  10-JAN-2002.
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05-MAR-1991
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                                                                                                                                                                                              Kazemi A,
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The sequence is that of a detection step primer for use in the identification of a mutation (G -> A) in the second mucleotide of codon 12 of the N-ras gene. It corresponds to nucleotides 15 to 34 on the N-ras gene and was synthesised on an Applied Biosystems 381A DNA synthesiser. It allows the accurate determn. of changes in the N-ras gene with such efficiency and ease that large numbers of samples can be screened. See also AAQ13677-Q13689. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detection of specific nucleotide variations - by primer extension using a detection step primer immediately adjacent the variable nucleotide.
                                                                                                                                         This nuclectide sequence is found in posn. 1388-1369 of HIV-1 Bru, 1421-1403 of HIV-1 Mal, 1388-1369 of HIV-Eli, 1706-1687 of HIV-2 ROD and 1670-1651 of SIV-MAC. It is the anti-sense strand of a primer pair used to amplify these HIV-1, HIV-2 and SIV viral sequences, esp. in conjunction with in vitro diagnosis of in-fection. It is useful for treating viral diseases, eg. AIDS. See also AAQ06905-08 and AAQ06910-54. (Updated on 09-JAN-2003 to add missing OS field.)
                                        New nucleotide sequences derived from genome of HIV-1, HIV-2 and SIV useful as primers for amplification of immuno-deficiency viruses in diagnosis and for raising antibodies in treatment of HIV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-ras gene codon 12 nucleotide variation detection step primer.
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                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 4 C; 5 G; 6 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 29; Page 61; 67pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                CIYIGCAIRGCIGCYIGA 18
                                                                                                                 Claim 2; Page 18; 24pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ13687 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                        Match 0.8%;
Local Similarity 72.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Soderlund H, Syvanen AC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ORIN ) ORION YHTYMAE OY
                                                                                                                                                                                                                                                                                                                                                           13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1991-281407/38.
          WPI; 1990-378039/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-FEB-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-FEB-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9113075-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ13687;
                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1195
                                                                                                                                                                                                                                                                                                                                           Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ1368'
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(revised)
(first entry)

90US-00482005 90US-00482005. 0.8%; Score 13.6; DB 1; Length 20;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This antisense oligonucleotide was designed to hybridise to the 3'-UTR thuman ICAM-1 mRNA. It was synthesised in the phosphorothicate form as none of the phosphodiester form-antisense oligonucleotides which were initially tested demonstrated inhibitory activity. Oligonucleotide #15 was found to be the most active of 16 potentially inhibitory anti-sense sequences. Its anti-sense activity was not shared by other oligonucleotides which hybridise to 3'-untranslated sequences. See e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New oligonucleotides hybridisable to cell adhesion modulators - for treatment and diagnosis of e.g. allograft rejection, cancer, AIDS etcand diagnosis of intercellular adhesion dysfunction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                          Antisense oligonucleotide #15 targetted to ICAM-1 3'-UTR (1952-1971).
               Gaps
                                                                                                                                                                                                                                  Intercellular adhesion molecule-1; inhibitor; phosphorothioate bond; triple helix; 3' untranslated region; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4; Indels
               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 13.6; DB 1;
Pred. No. 9.3e+02;
  80.0%; Pred. No. 9.3e+02; ive 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    K-ras codon 12 MTO-PCR set 1 primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAGAGTGGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 5; Page 43; 75pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                   Mira,
                                        229 AGTGGTGGTGGTGGCGCAG 248
                                                              20
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                                                                 1 Acresiseresiresascas
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                                                                                                                                                                                                                                                                                                                                                       91WO-US005209.
                                                                                                                                                                                                                                                                                                                                                                                 90US-00567286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ66488 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                  Bennett CF, Mirabelli CK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                 AAQ22643 standard; DNA; 20
                                                                                                                                                                                    08-JUL-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-096579/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
Best Local Similarity
Matches 16: Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-FEB-1995
                                                                                                                                                                                                                                                                                                                                                        23-JUL-1991;
                                                                                                                                                                                                                                                                                                                                                                                   14-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16,
                                                                                                                                                                                                                                                                                                                               05-MAR-1992.
                                                                                                                                                                                                                                                                                                      WO9203139-A.
                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ66488;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                           AAQ22643;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ66488,
                                                                                                        RESULT
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Length 20; 2; Indels

Score 13.6; DB 1; Pred. No. 9.3e+02; 3; Mismatches

1720

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02-SEP-1992;
21-JAN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9412617-A1
                                17-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
02-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-JUN-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                 20
                                                                                                                                                                                                                                                                                                                                                                                                            AAQ67992;
                                                                                                                                                                                                                                                                                                              226
                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                  RESULT 1199
                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                              AAQ67992,
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                                                                                                                                                                                                                                                                                                                                 셤
                                                                                                                                                                                                                               The sequences given in AAQG6488-90 are primers which were used in the method of the invention for the detection of a mutant oncogene. The method allows the detection and measuring a mutant oncogene contained in a sputum sample. PCR is performed by utilising the mutation position of the objective mutant oncogene as the complementary base to the 3' there by a bring a mixture of three primers which are different from teh the normal sequence at the 3' terminus. Another primer is used to hold the mutant oncogene together so that the mutant oncogene can be amplified position specifically and detected is pref. the K-ras gene and the mutation to be detected is pref. either codon 12, 13 or 61. This method allows detection of a mutation which is present only in trace amounts in the test sample
                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human intercellular adhesion molecule; ICAM-1; cell adhesion; modulation; inflammation; psoriasis; malignant melanoma; inflammatory bowel disease; antisense oligonucleotide; therapy; ss.
Polymerase chain reaction; primer; PCR; amplify; oncogene; K-ras; mutant;
                                                                                                                                                                            Detection of variant oncogene by PCR amplification - using the mutation site as the complementary base to the 3' end of a PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotide which targets human ICAM-1 3'-UTR.
                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/note= "in phosphorothioate form"
                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 3 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                       1311 GACATACAACTACCCCAAGT 1330
                                                                                                                                                                                                            Disclosure; Fig 1; 6pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 GAGCTCCAACTACCACAAGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ44522 standard; DNA; 20 BP.
                                                                                             92JP-00345280.
                                                                                                                 92JP-00345280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93WO-US008101
                                                                                                                                    (SAKA ) OTSUKA PHARM CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8
Best Local Similarity 80.0
Matches 16; Conservative
         detection; sputum; ss
                                                                                                                                                          WPI; 1994-230933/28.
                                                  JP06167492-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
misc_feature
                                                                                           30-NOV-1992;
                                                                                                                 30-NOV-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-AUG-1993;
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26-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-MAR-1994,
                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense oligonucleotides which target human ICAM-1 were synthesised in both the phosphodiester and phosphorothioate forms. The oligonucleotides are useful to treat diseases which are modulated by changes in intercellular adhesion molecules. This sequence corresponds to nucleotides 1952-1971 of the 3'- untranslated region of the human ICAM-1 coding sequence. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                            Oligo:nucleotide modulation of cell adhesion - used in the treatment of e.g. psoriasis, inflammatory bowel disease or malignant melanoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prevention and treatment of hepatitis - using recombinant replicable vaccinia viruses contg. hepatitis B virus surface and core antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence of PCR primer for modified HBV core antigen core delta 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Core antigen; recombinant replicable vaccinia virus; hepatitis; prevention; therapy; epitope; hepatitis B virus; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Souw PTS, Okeefe RW, Lewis T, Bernstine EG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 15; Page 51; 101pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAGAGTGGTGGTGGTGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ITBI-) INT BIOTECHNOLOGY LAB INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example; Page 84; 252pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAGAGGGAAGTGGTGGGGG 1
92US-00939855.
93US-00007997.
93US-00063167.
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AAQ67992 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                       Bennet CF, Mirabelli CK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 80.0
hes 16; Conservative
                                                                                                                                     (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleotide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1994-200247/24.
                                                                                                                                                                                                                                                                          WPI; 1994-100869/12.
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Gaps

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4; Indels

Length 20;

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8, in which 8 AAs are deleted, was used. A construct was made consisting of HBV MS antigen expressed from the modified p7.5 promoter and core delta 8 expressed from the p7.5 promoter. One primer used was AAQ67992 which hybridises from bases -230 to -211 relative to the ATG of core delta 8, upstream of the p7.5 promoter, and creases a HindIII site. A second primer (AAQ67993) hybridises to the opposite strand from bases +553 to +546 and generates a HindIII site. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ71022 and AAQ71023 are primers for the DNA sequence (AAQ79199) that codes for the rat mu-subtype opioid receptor (AAR65188). (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pure mu-type opioid receptor protein - and nucleic acid coding for it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus; Mu-subtype opioid receptor; MSOR; primer; ss.
                                                                                                                                            Sequence 20 BP; 3 A; 3 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                        Score 13.6; DB 1;
Pred. No. 9.3e+02;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer for the mu-subtype opioid receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Corbett MJ
                                                                                                                                                                                                                                     512 ACCTGGAGAGCTGACCCTC 531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          849 CCTGGACAAGGACCTGAAGC 868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 9; 39pp; English
                                                                                                                                                                                                                                                                   20 ACATCGAGAAGCTTACCCAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zysk JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCTGGACGAGAACTTCAAGC
                                                                                                                                                                            0.8%;
Similarity 80.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (AMCY ) AMERICAN CYANAMID CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94EP-0010196B
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                                                                                                                                                                                                                                                                                                                                              AAQ71023 standard; cDNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ71501 standard; cDNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                        16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eppler CM, Shieh H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1994-265963/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
19-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                             AAQ71023;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
                                                                                                                                                                        Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ71501
                                                                                                                                                                                                                                                                                                                  RESULT 1200
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                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ71501/
ID AAQ7
XX
AC AAQ7
   8888888888888
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Rapid detection of Brucella may be achieved by amplifying the omp2 gene locus of Errcella (which shows genetic variation correlating with established species designations) and hybridising the amplified sequence with a panel of DNA probes to identify a species of biovar of Brucella. The amplified sequence is preferably a sequence between nuclectides 2470 used for the consensus sequence described in AAQ1479. The method is used for the detection of Brucella infection in animals, particularly humans and cattle. This probe specifically hybridises to sequences from Brucella abortus biovar 1, Brucella abortus biovar 5, Brucella melitensis, Brucella neotomae and Brucella ovis which are amplified by the primers described in AAQ71496 and AAQ71497. The use of an array of probes (See AAQ71495-509) allows specific identification of the species of Brucella. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                          consensus; Brucella; identification; diagnosis; infection; biovar; s; disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                 Identification of Brucella species or biovars - by amplification of the Brucella omp2 gene locus and hybridisation with DNA probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Glutamate receptor; EAAS receptor; excitatory amino acid; CNS receptor; RNA editing; polymerase chain reaction; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                              Probe for identifying Brucella species.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Col 41; 50pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        177 CCGAGGCATAGACAAGACCA 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 CCGAGTCATAGGCAACAACA
                                                                                                                                                                                                                                                                                 90US-00527017.
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               (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                 (TEXA ) UNIV TEXAS A & M.
                                                                                                                                                                                                                                                                                                                                                   Ficht TA, Adams LG;
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1994-302203/37.
                                                                                               omp2; consensus; cattle; disease;
               25-MAR-2003
02-MAY-1995
                                                                                                                                                                                                                                               06-NOV-1992;
                                                                                                                                                                                                                                                                               22-MAY-1990;
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                                                                                                                                                                                                                 20-SEP-1994.
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                                                                                                                                                                                 US5348857-A
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                                                                                                                                               Synthetic.
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Gaps

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226 GAGAGTGGTGGTGGCGG 245
                                           Claim 2; Page 35; 57pp; English
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94US-00214582.
94US-00215088.
94US-00227369.
94US-00251938.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-MAY-1996 (first entry)
                                                                                                                                                                                                                                                                                                         Local Similarity 80.0 tes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1995-344401/44.
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18-MAR-1994;
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AAQ99937/c
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/*tag= a
/*tag= a
/note= "at least one (and preferably all) of the backbone
subunits are composed of amide units, so that the
subunits are consists of the nucleobases attached covalently
to a polyamide backbone"
                                                                                                                                                                PCR primers (AAQ91246-50) were used to amplify human glutamate receptor EAAS genomic DNA and cDNA. Examination of the PCR products showed that the cDNA sequence differed from the genomic Sequence at 2 places in the transmembrane domain-coding region, resulting in S310A and R5220 substitutions. These variations were attributed to RNA editing involving T to G and G and A substitutions. Similar RNA editing was found for EAA3 (see also AAQ91231) and EAA4 (see also AAQ91232) genes
                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide nucleic acid; PNA; intercellular adhesion molecule; ICAM-1; endothelial leukocyte; BLAM-1; vascular; VCAM-1; antiinflammatory; anticancer; antimetastatic; anti-AIDS; anti-rhinoviral; ss.
                                                                                                               Identification of human CNS receptor ligand - and identification of agents that modulate editing of human CNS receptors.
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                                                                                                                                                                                                                                                                         0.8%; Score 13.6; DB 1; Length 20;
00.0%; Pred. No. 9.3e+02;
.ve 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide Nucleic acid oligomer targetting ICAM-1 3'-UTR.
                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 4 C; 9 G; 4 T; 0 U; 0 Other;
                                                  (ALLX ) ALLELIX BIOPHARMACEUTICALS INC.
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                                                                                                                                                                                                                                                                                                                    1211 CGGGCTCCACGGTGGAGGAA 1230
                                                                                                                                              Example 9; Page 35; 59pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                          AAT01753 standard; DNA; 20 BP.
          94WO-CA000705.
                              93US-00172188
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Best Local Similarity 80.05
Matches 16; Conservative
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                                                                                           WPI; 1995-240670/31.
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                                                                       Kamboj R, Nutt S;
                             23-DEC-1993;
           21-DEC-1994;
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New oligomers are claimed which (A) have at least one peptide nucleic acid (PNA) subunit and (B) have a sequence hybridisable to AUG region.

Coding region, 5'-untranslated region or 3'-untranslated region of ICAM-1 or ELAM-1, or hybridisable to AUG region, coding region, 5'- untranslated region of UCAM-1 or ELAM-1, or hybridisable to AUG region, coding region, exon/intron junction region or 3'-untranslated region of VCAM-1.

The PNAs can be used to target RNA and single stranded DNA (SDNA) to produce antisense-type gene regulation moieties. Hence they may be used therepoetically for modulating cellular adhesion and thus as an antimetastatic agents, anticancer agents, anticancer agents, anticancer agents, antiching callular adhesion and thus as an antimetastatic agents, anticancer agents. They may also be useful as antimetastatic agents, anticancer specific mRNAs. PNA oligomers have high affinity for complementary single stranded DNA. They are also able to form triple helices in which a first PNA strand binds with the resulting double helix or with the first PNA strand. The PNAs possess no significant charge and are water soluble, which facilitates cellular uptake. Further, since they contain amides of non-biological amino acids, they are bloestable and resistant consymatic degradation by proteases. The present sequence targets human cut intercellular adhesion molecule-1 (ICAM-1) 3' untranslated region
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New peptide nucleic acid oligomers hybridising to adhesion molecule genes - are stable anti:sense cpds. of high affinity, partic. for treating inflammation, viral infection, cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wild-type multiple tumour suppressor (MTS) gene and mutant sequences - useful in diagnosis, prognosis and therapy of human cancer, e.g. melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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80.0%; Pred. No. 9.38+02;
ive 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
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                                                     The cDNA sequences encoding several multiple tumour suppressor (WTS) polypeptides have been isolated and sequenced, using various sequencing and amplification primers. AAG99936-40 are oligomuclectides used to amplify CDNA encoding mouse WTSIE1-beta to allow comparison of the human and murine sequences. WTS polypeptide-encoding cDNAs and mutants of themen are useful for the diagnosis or prognosis of human cancer. Germ-line mutations of MTS cDNAs can be used for diagnosing predisposition to melanoma, leukaemia, astrocytoma, glioblastoma, lymphoma, glioma, Hodgkin's lymphoma, CLL and cancers of the pancreas, thyroid, ovary, Hodgkin's lymphoma, testis, kidney, stomach and rectum. The wild-type gene is useful for gene therapy and MTS polypeptides may also be used for protein replacement therapy. Also the polypeptides or cells contg. an altered MTS gene are useful for screening for potential cancer therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQB1115 is a peptide nucleic acid (PNA), which binds a target sequence. The binding of the PNA prevents the transcription of the target sequence
                                                                                                                                                                                                                                                                                                                                                Gaps
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binding to ssDNA, dsDNA or RNA for use in therapy, diagnosis and
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                                                                                                                                                                                                                                                                                                            0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels
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/note= "covalently bound Lys-NH2 group"
                                                                                                                                                                                                                                                                                Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        outnardt O, Egholm M, Nielsen PE, Mollegaard NE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                              Example 12; Page 71; 156pp; English
                                                                                                                                                                                                                                                                                                                                                                              505 GAGGCTACCTGGAGAAGCT 524
                                                                                                                                                                                                                                                                                                                                                                                                GAAGGCTTCCTGGACACGCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
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(first entry)
                                                                                                                                                                                                                                                                                                                                                16; Conservative
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                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      prophylaxis; ss
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modified_base
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28-SEP-1995
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Matches
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AAQ81115/
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by RNA polymerase. The ability of the PNA to arrest transcription makes it useful in gene therapy, and in diagnostic and prophylatic methods. (Updated on 25-MAR-2003 to correct PN field.)
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binding to ssDNA, dsDNA or RNA for use in therapy, diagnosis and
prophylaxis.
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                                                                           0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
ative 0; Mismatches 4; Indels
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                                                    Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
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/note= "amidated"
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(first entry)
                                                                                Query Match
Best Local Similarity 80.0
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                            Peptide nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                     prophylaxis; ss
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Mollegaard NE;
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modified_base
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28-SEP-1995
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RESULT 1207

AAQ80945

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antisense; analogue; non-terminal pyrimidine; phosphorothioate; backbone; treatment; HIV; human immunodeficiency virus; HSV; herpes simplex virus; cancer; integrin; cell adhesion receptor; infection; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                           An individual can be diagnosed as having a predisposition to cancer by detecting an alteration in the wild type multiple tumour suppressor (MTS) gene, using gene probes which hybridise to the MTS1 gene (amplified using the PCR primers AAT00729-31). The above assay can also be used in the diagnosis and prognosis of melanoma, lymphoma, leukaemia and pancreas, breast and thyroid cancers, etc
                                                                                                                                                                                                                                                                                                                                   Detecting polymorphism associated with cancer pre:disposition - also DNA, vectors and host cells e.g. for gene or protein replacement therapy and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
rative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human ICAM modified antisense oligonucleotide.
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                                                                                                                                                                                                                                                                   Kamb
                                                                                                                                                                                                                                                                                                                                                                                                                Example 12; Page 71; 148pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         505 GAGGCTACCTGGAGAGCT 524
                                                                                                                                                                                                                                                                   Cannon-Albright LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 GAAGGCTTCCTGGACACGCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВЪ.
                                                                                                94US-00214582.
94US-00215086.
94US-00215087.
94US-00227369.
94US-00251938.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94EP-00117513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93DE-04338704
                                                                 95WO-US003537
                                                                                                                                                                                                               (UTAH ) UNIV UTAH RES FOUND (MYRI-) MYRIAD GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ88741 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 80.0 nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nuclease resistance; ss.
                                                                                                                                                                                                                                                                                                      WPI; 1995-344626/44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (FARH ) HOECHST AG.
                                                                                                                                                                                                                                                                                                                                                                                 drug screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-NOV-1993;
                                                                                                                                                                                                                                                                     Skolnick MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-FEB-1996
WO9525813-A1.
                                                                   17-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP653439-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-MAY-1995.
                                                                                                                                       18-MAR-1994;
14-APR-1994;
                                                                                                                                                                              01-JUN-1994;
                                                                                                      18-MAR-1994;
                                28-SEP-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ88741;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 12
AAQ88741/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A probe of approximately 1 kb recognising beta-giardin genomic DNA was generated by PCR with the oligonucleotides AAQ80942 and AAQ80943. A random probe was generated as a 1.5 kb product with the primers AAQ80944 and AAQ80945. The probes were used in the identification of cosmids from the beta-giardin genomic region in a diardia lamblia 20-genome equivalent cosmid library. This was part of a novel method for sequence-sampled mapping of complex genomes. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ል
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequencing complex genomes, present as fragments in a cosmid library - k sequencing end-specific nucleotides of each clone then correlating with spatial relationship of cosmid, esp. for mammalian chromosomes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Multiple tumour suppressor; MTS1; cancer; diagnosis; assay; predisposition; melanoma; leukaemia; lymphoma; prognosis; pancreas; breast; thyroid; P16 specific; reverse PCR primer; ss.
                                                                                                                                                                                                               sequence sampled mapping; genomic analysis; complex genome mapping; cosmid library; Giardia lamblia; ss.
                                                                                                                                                                                  PCR primer to generate a random probe for screening complex genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Multiple tumour suppressor 1 gene P16 specific reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
.ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 5 A; 11 C; 1 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SALK ) SALK INST BIOLOGICAL STUDIES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 43; 128pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     230 GTGGTGGTGGTGGCGCAGT 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 GAGGTGGTGTGTCAGGAGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                    AAQ80945 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                     94WO-US006810.
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93US-00117952.
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                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 80.0
Matches 16; Conservative
                                                                                                                             (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Evans GA, Smith MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1995-036508/05.
                                                                                                                                                                                                                                                                                                                                                                                   15-JUN-1994;
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07-SEP-1993;
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                                                                                                                                                                                                                                                                                                             WO9429486-A1
                                                                                                                             25-MAR-2003
24-AUG-1995
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                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                          AAQ80945;
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Gape ö Winkler I;

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Helaberg

Kretzschmar G,

Mag M,

Peyman A, Uhlmann E, WPI; 1995-180677/24

Synthetic

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The antisense oligonucleotide (ON) shown is a derivative of an equivalent wild type Human ICAM ON, in which at least one, esp. 2-10, non-terminal pyrimidine nucleotide(8) is/are modified. The modification may be: (a) replacement of a phosphodiseter linkage by: a phosphoro-thioate (PS), dithioate, aramidate; borano-, alkyl-, aralkyl-phosphore: 2,2.2-trichloro-1, Idimethyl-, alkyl- or aryl- phosphorate linkage; or (3'-thio) formacetal, methylhydroxylamine, oxime, methylenge; to replacement of a sugar dimethylene sulphone or silyl linkage; (b) replacement of a sugar phosphorae backbone by a "morphholinonucleoside" oligomer; (c) replacement of beta-D-2-deoxyribose by another sugar or carbocyclic, open-chain or bicyclic sugar analogue, or (c) replacement of the natural nucleoside base by an analogue, e.g. 5-hydroxymethyl-uridine. The 5' and/or 3' terminus may also be modified with a lipophilic gp, eg, a farnesyl. The modifications increase nuclease resistance and thus improve stability and
                 New anti:sense oligo:nucleotide analogues - with modified non-terminal pyrimidine nucleotide units, useful for treating viral infections,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene signature; messenger RNA; mRNA; relative abundance; frequency; human; cloning; mapping; non-biased library; diagnosis; detection; cell typing; abnormal cell function; primer; PCR; amplification; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human gene signature HUMGS00995-derived anti-sense primer.
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 cacaccaactccreccc
                                                                                      Claim 1; Page 31; 36pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT41336 standard, DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1995-206931/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MATS/) MATSUBARA K. (OKUB/) OKUBO K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-NOV-1994;
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                                                      cancer, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-DEC-1996
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0; Gaps

Single-stranded DNA for identifying gene signatures - isolated from 3'-directed human cDNA library that reflects relative abundance of corresp. mRNA in specific human tissues.

Primers T41001-T41382 are derived from novel human gene signature (GS)

Example 7; Fig 10; 2245pp; Japanese.

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sequences which did not match with sequences deposited in Genbank release 76. The GS sequences (T19001-T26837) were obtained from 3'-directed cDNA libraries prepared from various human tissues, synthesis of cDNA was initiated from the 3'-end of mRNA by using poly(T) as the sole primer. Bach library is constructed so as to reflect accurately the relative abundance of different mRNAs in the particular tissue from which it was determined (sp. using primers and probes derived from the GS sequences) as a means of diagnosing abnormal cell function or for recognising different cell types. The primers $71335-6 amplify clone pm0268 which comprises the GS HUMSS000955 (T19995). This amplification reaction gave prod. indistinguisable from the same PCR using mouse or Chinese hamster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A sense oligonucleotide S50 (AAQ99516) corresp. to nucleotides 50-69 in the human Fas ligand coding sequence given in AAT03498 was synthesised with phosphororhioate linkages. The complementary, antisense oligonucleotide A69 (AAQ99517) was also synthesised. The effects on Fas ligand-mediated apoptosis of A69 and S50 were analysed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide which binds to Fas antigen, and antibody reactive with it - for treatment and diagnosis of viral or auto:immune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fas ligand, Tumour Necrosis factor family; apoptosis; cell death; Fas cell surface antigen; human; Fas-L; phosphorothioate; antisense oligonucleotide; inhibition; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Fas ligand phosphorothicate antisense oligonuclectide A69.
                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                        0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                             Sequence 20 BP; 5 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 5 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nakamura N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 20; Page 111; 300pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                    339 GGACTTGAAGATGGGGTCTG 358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suda T, Takahashi T,
                                                                                                                                                                                                                                                                                                                                                                                                     20 GGTATAAAGATGGGGTCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ99517 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-FEB-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                   16; Conservative
                                                                                                                                                                                                                          ovary DNA as a template
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1995-194031/25.
                                                                                                                                                                                                                                                                                                             Local Similarity
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08-JUL-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ99517;
                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1211
                                                                                                                                                                                                                                                                                                                                   Matches
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RESULT 1212

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New oligo:nucleotide(s) contg. 8-aza:purine base - useful as therapeutic and diagnostic agents with more stable hybridisation to target nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT44425-54 are antisense oligonucleotides which have at least one 8-azapurine base. The presence of an 8-azapurine base results in significantly stronger complexing when hybridising to target nucleic acids. The present sequence is against the intracellular adhesion molecule (ICAM) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense therapy; guanosine; intercellular adhesion molecule; ICAM; nuclease resistance; stability; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                               8-azapurine, modification; stronger complex, inhibition; intracellular adhesion molecule; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ICAM antisense component of capped oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                Antisense oligonucleotide against ICAM gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 44; 51pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 GAGAGGGAAGTGGTGGGGG 1
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AAT44250 standard; DNA; 20 BP.
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              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                WPI; 1995-375165/49
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                                                                                                                                                                                                                                                                                                           (FARH ) HOECHST AG.
                                                                                                                                                                                                                                                                                                                                              Seela F, Lampe S;
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                                                                                                                                                                                                                                                                          02-MAY-1994;
                27-JAN-1997
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                                                                                                                                                                                                         08-NOV-1995
                                                                                                                                                                      EP680969-A2
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                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peyman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT44250;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT44250/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A sense oligonucleotide S50 (AAQ99516) corresp. to nucleotides 50-69 in the human Fas ligand coding sequence given in AAT03498 was synthesised with phosphorothioate linkages. The complementary, antisense oligonucleotide A69 (AAQ99517) was also synthesised. The effects on Fas ligand-mediated apoptosis of A69 and S50 were analysed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide which binds to Fas antigen, and antibody reactive with it - for treatment and diagnosis of viral or auto:immune diseases.
                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                         Fas ligand; Tumour Necrosis factor family; apoptosis; cell death;
Fas cell surface antigen; human; Fas-L; phosphorothicate;
sense oligonucleotide; inhibition; ss.
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                                                                                                                                                                                                                                                                                                           Human Fas ligand phosphorothioate sense oligonucleotide S50.
           Ouery Match

0.8%; Score 13.6; DB 1; Length 20;
Best Local Similarity 80.0%; Pred. No. 9.3e+02;
Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 5 A; 8 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Takahashi T, Nakamura N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 20; Page 111; 300pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      483 ACCAGCTGACATCCGGCTGC 502
                                                                                   483 ACCAGCTGACATCCGGCTGC 502
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                                                                                                                  20 Accaderecearecades 1
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93JP-00342526.
94JP-00074344.
94JP-00180955.
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                                                                                                                                                                                                         AAQ99516 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94JP-00278378
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                                                                                                                                                                                                                                                                          28-FEB-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-194031/25.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-DEC-1993;
18-MAR-1994;
08-JUL-1994;
07-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9513293-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                         AAQ99516;
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RESULT 1213

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AAT44449,

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Gaps

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This sequence represents an inhibitor of ICAM, and is an example of an oligonucleotide analogue of the invention. The oligonuclectide analogues of the invention are used as inhibitors of gene expression (antisense oligonuclectides, ribozymes, sense oligonuclectides and triplex-forming oligonuclectides), as probes for the detection of nucleic acids, and as auxiliaries in molecular biology. As gene expression inhibitors they may be used for treating viral infections (especially where the virus is HSV-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene expression inhibitor; probe; nucleic acid detection; growth factor; viral infection; therapy; HSV-1; cancer; restenosis; integrin; cell-cell adhesion receptor; ICAM; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New phosphono:mono:ester oligo:nucleotide analogues - inhibitors of gene expression for treating viral infections, cancer, restenosis, etc.
                                                Oligo:nucleotide(s) with series of G residués at at least one end have increased stability against nuclease and cell penetration, - are partic. anti:sense sequences for treating and diagnosing cancer, viral diseases
                                                                                                                                                                           Ten- to 40-mer oligonucleotides which have a cap of 1-10 (esp. 4) G residues on at least one end are provided; if caps are present at both ends, they can be of the same or different lengths. A cap sequence increases nuclease resistance of the oligonucleotide and also increases cell penetration. The present sequence is that of a preferred oligonucleotide, directed against an intercellular adhesion molecule sequence, which can be capped for use in anticancer therapy
                                                                                                                                                                                                                                                                                                                                                             o.8%; Score 13.6; DB 1; Length 20; Similarity 80.0%; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 42; 129pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Breipohl G,
                                                                                                                                                                                                                                                                                                                                                                                                                                   226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dadadedeaadrecreeed 1
                                                                                                                                          Claim 3; Page 13; 15pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96AU-00048028
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95DE-01043865.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX33922 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8
Best Local Similarity 80.0
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ICAM expression inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peyman A, Uhlmann E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-455932/46.
                 WPI; 1996-355223/36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-MAR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-NOV-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX33922;
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AAX33922/c
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Wallmeier H;

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1, HSV-2, an influenza virus, VSV, hepatitis B or papilloma virus), cancer, restenosis, medical conditions mediated by integrins or cell-cell adhesion receptors, and medical conditions induced by growth factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Transgenic animals contg. an immortalising gene and intestinal neoplasia-
associated gene - useful in studies of e.g. carcino-genesis, that may be
induced by e.g. viruses, various gene or mutagens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; PCR; polymerase chain reaction; amplification; detection; Min; multiple intestinal neoplasia; cancer; colon; SV40 large T gene; transgenic mouse; immortalised cell line; ss.
                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer for Min mutant allele 280 bp PCR product amplification.
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80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                            Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 9.3e+02; les 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 20 BP; 4 A; 0 C; 6 G; 10 T; 0 U; 0 Other;
                                                                                          Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1309 AAGACATACAACTACCCCAA 1328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 4; Page 17; 50pp; English.
                                                                                                                                                                                                         226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 AACACATACAACTTCACTAA 1
                                                                                                                                                                                                                                          20 dadadeceaadrecreece 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (LUDW-) LUDWIG INST CANCER RES.
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                                                                                                                                                                                                                                                                                                                                     AAT15587 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Whitehead RH, Joseph JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 80.0 nes 16; Conservative
                                                       (especially TNF-alpha)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-068869/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                              Query Match
Best Local S:
Matches 16
                                                                                                                                                                                                                                                                                                                                                                          AAT15587;
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                                                                                                                                                                                                                                                                                                   RESULT 1216
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Gaps ö

4; Indels

20 GAGAGGGAAGTGGTGGGGG 1

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AAX24204

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AAT30211-T30233, AAT33058-T33112 and AAT36667-T36684 represent antieense oligonuclectides of the invention. These sequences target regions of the coding sequences for human intercellular adhesion molecule-1 (ICAM-1), endothelial leukocyte adhesion molecule-1 (ELAM-1, also known as Eschectin), or vascular cell adhesion molecule-1 (VCAM-1). This sequence targets the 3' untranslated region (uncleotides 1952-1971) of ICAM-1. ICAM-1, ELAM-1, and VCAM-1 represent three of the five cell adhesion molecules involved in the adherence of white blood cells to vascular endothelium. These sequences can be used in a composition for treating callograft rejection. The composition contains one of these sequences in combination with an immunosuppressive agent. The immunosuppressive agent cused in the compositions is brequinar, rapamycin, anti-lymphocyte serum, a monoclonal antibody against LFA-1 or an antisense oligonucleotide. The such as cardiac or renal allograft rejection. By using these compositions, allograft rejection. By using these compositions, allograft rejection. By using these
                                                                                                                                                                      Antisense oligonucleotide; human; intracellular adhesion molecule-1; ICAM-1; endothelial leukocyte adhesion molecule-1; ELAM-1; E-selectin; vascular cell adhesion molecule-1; VCAM-1; white blood cell; brequinar; vascular endothelium; lograft rejection; immunosuppression; rapamycin; anti-lymphocyte serum; monoclonal antibody; cardiac allograft; therapy; renal allograft rejection; donor-specific transplant tolerance; IFA-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligo:nucleotide targetted to a nucleic acid sequence encoding ICAM-1, ELAM-1 or VCAM-1 - useful for treating or preventing allo:graft rejection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
ative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= a
/note= "phosphorothicate backbone"
                                                                                                                                          Antisense oligonucleotide ISIS 1939.
                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 49; 92pp; English.
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                                 AAT30227 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95WO-US015536.
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(TEXA ) UNIV TEXAS SYSTEM.
                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                     /*tag=
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                                                                                                                                                                                                                                                                                                                                                                      Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9615780-A1
                                                                                                       20-JAN-1997
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                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                      AAT30227;
RESULT 1217
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This invention describes novel phosphonomonoester oligonuclectide analogues which act as inhibitors of gene expression (as sense/antisense, ribozyme or triplex-forming molecules), useful as diagnostic agains (i.e. probes for detecting nucleic acid) or for treatment of diseases caused by viruses, influenced by integrins or cell-cell adhesion receptors, induced by factors such as TMF-alpha, or cancer or restencies. The products of the invention satisfy the requirements of good in-vivo stability; ability to cross cellular and nuclear membranes, and specific binding to target nucleic acid better than known oligonucleotides
                                                                                                                        Phosphonomoncester analogue; inhibitor; antisense; cancer; restenosis; ribozyme; diagnostic agent; detection; treatment; disease; virus; integrin; cell-cell adhesion receptor; TNF-alpha; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense; anti-proliferative; tumour; cancer; raf; oncogene; phosphorothicate; 2' sugar modification; psoriasis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            New oligo:nucleotide analogues contg. phospho:mono:ester bridges -
therapeutic inhibition of gene expression, e.g. in cancer or viral
infection, with good specificity and in vivo stability.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human c-raf kinase coding region antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                Wallmeier
                                                                                               Phosphonomonoester oligonucleotide analogue 21
                                                                                                                                                                                                                                                                                                                                                                                                Breipohl G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 23; 36pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 GAGAGGGAAGTGGTGGGGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                 Uhlmann E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
AAX24204 standard; DNA; 20
                                                               01-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-425893/43.
                                                                                                                                                                                                                                                                                                                                                                   (FARH ) HOECHST AG.
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                                                                                                                                                                                                                                  DE19508923-A1.
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                                                                                                                                                                                                                                                                                                    13-MAR-1995;
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                                                                                                                                                                                                                                                                    19-SEP-1996.
                                                                                                                                                                                                  Synthetic.
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                                   AAX24204;
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Query Match
Best Local Similarity 80.0
Matches 16; Conservative

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Gaps

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95WO-US007111 94US-00250856

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Oligo:nucleotide(s) targetted to nucleic acids encoding human raf capable of inhibiting raf expression, used in treatment of hyper:proliferative disorders.
                                                                                                                                                                                                                                                                                                            Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 9.3e+02; les 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human potassium channel gene subregion E antisense primer.
                                                                                                                                                                                                                                                                                            Sequence 20 BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYCH-) UNIV CHIBA SUSUMU SEINO INOHANA SHUKUSHA.
                                                                                                                                                                                     Disclosure, Page 15, 65pp; English.
                                                                                                                                                                                                                                                                                                                                                   1186 ATGGCCACAGGCCGTCCCCT 1205
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                                                                                                                                                                                                                                                                                                                                                                                                               AAT61877 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JCR PHARM CO LID
                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seino S, Inagaki N;
                                                                                                                Monia BP, Boggs RT;
                                                                                                                                 WPI; 1996-030518/03.
                                                                             31-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-1996;
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                                                            31-MAY-1995;
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                         WO9532987-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP764721-A1
                                         37-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
        Synthetic.
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Best Local Si
Matches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT61877;
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                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1220
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В В

(first entry)

96EP-00114885 95JP-00264943

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= a
//note= ackbone selected from: phosphorothioate;
dithicate; methylphosphorate; phosphodiester; morpholino
backbone; polyamide backbone; and any combination of
these backbone types; the backbone may be modified to
incorporate a ribozyme structure, or a pendant group"
                                          Human and mouse pancreatic ATP sensitive potassium channel proteins - fo:
diagnosis, therapy and research into potassium channel related diseases,
e.g. diabetes.
                                                                                                                                                             PCR primers (AAT61868-79) were designed to amplify subregions A-F of the human pancreatic ATP sensitive potassium channel beta-IR gene (see also AAT61866). The antisense primer (AAT61877) for subregion B (494 bp) corresponds to nucleotides 999-1019 of the gene. Cy5-labelled primers were used in the PCR-SSCP analysis of genomic DNA collected from 20 healthy Japanese subjects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents a novel oligonucleotide 3(3B) MRP that
                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human multidrug resistance-1; MDR-1; inhibition; aptameric;
human multidrug resistance-associated protein; antisense; cytotoxic;
chemotherapeutic; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                     0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred, No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                      Seguence 20 BP; 3 A; 4 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Complementary human MRP oligonucleotide 3(3B) MRP
                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 17; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                      885 TGGGAACATCATCAACATGC 904
                                                                                                                                Example 3; Fig 6; 16pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 reschackechrehaerse 1
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                                                                                                                                                                                                                                                                                                                                                               80.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT48972 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-SEP-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                               Local Similarity 80.0
nes 16; Conservative
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/*tag=
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                  WPI; 1997-181836/17.
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misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-DEC-1996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT48972;
                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT27481-T27507 are human c-raf kinase antisense oligonucleotides used for the inhibition of raf expression. The oligonucleotides (ONs) are trageted to either codding region, start or stop signals or 5' or 3' untranslated region (UTR) mNAN encoding human c-raf. The ONs may be phosphorothioate linked and may contain modifications at the 2' position of the sugar molety. ONs are pref. complementary to either 3' or 5' UTRs, phosphorothioate linked and contain 2''O-alkyl sugar modifications. The ONs are used to inhibit expression of human raf in partic. in conditions associated with hyperproliferation e.g. cancer, restenosis, and psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATP-sensitive potassium channel protein; beta-IR; diabetes; pancreas; beta-cell; diagnosis; gene therapy; primer; PCR; polymerase chain reaction; single strand conformation polymorphism; SSCP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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20 GAAGGCTTCCTGGACACGCT 1

AAT98014 standard; DNA; 20

(revised)

08-SEP-1998

25-MAR-2003

AAT98014;

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specifically hybridises in a human cell with a complementary sequence of human multidrug resistance-associated protein (RRP) gene. Hybridisation causes inhibition of expression of the multidrug resistance phenotype by the cell, due to the oligomucleotide having an aprameric inhibitory effect as well as an antisense inhibitory effect. The oligomucleotide is administered to cancer patients to prevent development of the multidrug resistant phenotype. When co- administered with chemotherapeutic agents, the oligomucleotide is useful for potentiating alimination of multidrug resistant tumour cells from home marrow or peripheral stem cell grafts. Also, the oligomucleotide can be used as an immunosuppressive agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer; polymerase chain reaction; PCR; amplification; P16; promoter; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human mutant multiple tumour suppressor gene sequences - for production of recombinant mutant polypeptide (s).
                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is primer for the PCR amplification of the P16 promoter. (Updated on 25-MAR-2003 to correct PF field.)
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                                                                                                                                                                                                                      Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                           Sequence 20 BP; 1 A; 5 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Skolnick MH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pl6 promoter specific reverse primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 12; Col 83-84; 72pp; English
                                                                                                                                                                                                                                                                                          733 GCACCCTGCACCGCCATCCG 752
                                                                                                                                                                                                                                                                                                                           20 GCAGCAGCCACCGCCATCCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94US-00214582.
94US-00215086.
94US-00215087.
94US-00227369.
94US-00251938.
95WO-US003537.
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                                                                                                                                                                                                                                                                                                                                                                                                              AAT72304 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
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10-SEP-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT72304;
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comprising conserved sequences of human immunodeficiency virus and simian
immunodeficiency virus genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The oligonucleotides AAT98010-T98059 are useful as primers for nucleic acid amplification of conserved sequences of the gag, vpr. pol or vpu genes of the HIV-1 strains Bru, Mal, Bil, HIV-2 ROD or sinian immunodeficiency virus (SIV) MAC or the nef2, vif2 or vpx genes of HIV-2 ROD and SIV MAC. This primer is targetted to sequences in the gag gene of the viral strains. The sequence are therefore used to detect HIV-1, HIV-2 or SIV infections. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                   Primer; PCR; amplification; gag; vpr; pol; vpu; HIV-1; HIV-2; SIV; nef2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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0
                                                                                       Human or simian immunodeficiency virus detection primer MMy4B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer #35 for cystic fibrosis transmembrane regulator gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.6; DB 1; Length 20; 72.2%; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A; 4 C; 5 G; 6 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                            (INSP ) INST PASTEUR.
(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 4; Page 18; 23pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 cryrecaracciecyrea 18
                                                                                                                                                                                              Simian immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT47409 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                      97EP-00110543.
                                                                                                                                                                                                                                                                                                                  89FR-00007354.
                                                                                                                                                                                                                                                                                                                                  89FR-00012371.
90EP-00401520.
                                                                                                                                                                               immunodeficiency virus
                                                          (first entry)
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                                                                                                                                   vif2; vpx; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                        Moncany M, Montagnier L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-538622/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                      05-JUN-1990;
                                                                                                                                                                                                                                                                                                                   02-JUN-1989;
                                                                                                                                                                                                                                                                                                                                  20-SEP-1989;
                                                                                                                                                                                                                                                                                                                                                 05-JUN-1990;
                                                                                                                                                                                                                                                       12-NOV-1997.
                                                                                                                                                                                                                           EP806484-A2
                                                                                                                                                                 Synthetic.
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Gaps

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505 GAGGGCTACCTGGAGAAGCT 524

schultz621-3.rng

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transmembrane regulator (CFTR) gene. These sequences can be used as half transmembrane regulator (CFTR) gene. These sequences can be used as half of the chimeric primers of the invention. The primers are used for amplification of a target DNA sequence, and can be used in a multiplex per amplification. The primers have the sequence 5'-XY-3', where X is a sequence that does not hybridise to the target sequence (such as APT47344 and Y is a sequence contained within or flanking the target requence (such as this sequence). During early cycles of amplification, products are synthesised that contain the chimeric primers on either end. The primers then serve as high stringency recognition sequences for subsequent rounds of amplification. As a result, the annealing efficiency of different primers and their targets in a multiplex amplification or reaction is normalised, thereby reducing preferential amplification of certain targets. The chimeric primer comprise a 5' universal domain and a 3' target-specific domain. They are used for the simultaneous pcR amplification of multiple DNA targets in a sample. The primer containing AAT4744 is particularly useful in high-throughput genetic screening for mutations in genes like the CFTR, the Wilms Tumour, and the beta-
                  PCR; primer; amplify; polymerase chain reaction; bacteriophage; M13mp18; cystic fibrosis transmembrane conductance regulator gene; multiplex PCR; chimeric primer; genetic screening; mutation detection; CFTR; Wilms Tumour gene; beta-thalassaemia gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Universal primer used for multiplex DNA amplification - allows simultaneous amplification of multiple DNA target sequences for high through-put genetic screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Fig 1b; 38pp; English
                                                                                                                                                                                                                                                                                               95US-00474450.
                                                                                                                                                                                                                                                        96WO-US009637.
                                                                                                                                                                                                                                                                                                                                        (GENZ ) GENZYME CORP
                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1997-052372/05.
                                                                                                                                                                    WO9641012-A1
                                                                                                                                                                                                                                                                                                 1995;
                                                                                                                                                                                                               19-DEC-1996
                                                                                                                           Synthetic.
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Gaps . 0 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels Sequence 20 BP; 6 A; 5 C; 4 G; 5 T; 0 U; 0 Other; 16; Conservative Local Similarity Query Match ઠે

1223 TGGAGGAACAGCTACACTTC 1242 20 rádadozákokorakorrrro

g

AAT94038 standard; cDNA; 20 (first entry) (revised) 01-APR-1998 25-MAR-2003 AAT94038; RESULT 1225

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Canalicular multispecific organic anion transporter protein; cMOAT protein; ATP-binding cassette transporter family; ABC transporter; Forward PCR primer used to amplify a 241 bp fragmnet of CMOAT CDNA

canaldualar multispecific organic anion transporter (CMOAT) protein CDNA.
The PCR product was cloned, and subsequently used in a RNAse protection assay. CMOAT is a new member of the ATP-binding cassette (ABC) transporter family. The ATP dependent CMOAT transporter system mediates transporter family. The ATP dependent CMOAT transporter system mediates comported family. The ATP dependent CMOAT transporter system mediates transporter family. The ATP dependent CMOAT protein. The mucleic acids are used to provide cells with CMOAT protein. The mucleic acids are used to provide cells with CMOAT protein activity, CMOAT protein.

Care used to provide cells with CMOAT protein activity, CMOAT protein activity in cells can be enhanced by increasing the level of glutathione, glucuromide and/or sulphate. Antisense constructs, especially derived from another multidary resistance (MDA) related protein, e.g. MDAP-1, to the mucleic acids and vectors can be used to decrease the level of cMOAT in a cell. The mucleic acids and proteins can be used especially in change of Dubin-Ohnson disease, Rotor disease or another disease involving cMOAT. The CMOAT gene may also be used as a selectable marker gene. (Updated on 25-MAR-2003 to correct PI field.) hepatobiliary excretion; multidrug resistance-associated protein; CMOAT protein activity; multidrug resistance-related protein; MDR-1; Dubin-Johnson disease; Rotor disease; PCR primer; ss. DNA encoding human and rat canalicular multispecific organic anion transporter proteins - useful for diagnosis and treatment of Dubin-Johnson disease and Rotor disease. ä Borst P, Sequence 20 BP; 4 A; 6 C; 3 G; 7 T; 0 U; 0 Other; Bosma PJ, (INTR-) INTROGENE BV. (MEDI-) ACAD MEDISCH CENT AMSTERDAM. (HETN-) HET NEDERLANDS KANKER INST. Example 6, Page 29; 106pp; English. Oude Elferink RPJ, Paulusma CC, Kool M_I 97WO-NL000079. 96EP-00200460 WPI; 1997-435163/40. Homo sapiens. WO9731111-A2 21-FEB-1997; 28-AUG-1997. Synthetic.

Nucleotide sequence of P16 specific reverse PCR primer. CTTCATCTTCCGTATCTTAG 1258 creccrcrrcagaarcrrag 20 BP AAV53844 standard; DNA; 20 04-DEC-1998 (first entry) AAV53844; 1239 셤

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Gaps

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0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels

16; Conservative

Matches

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Local Similarity

Query Match

Multiple tumour suppressor; MTS; human; cancer; hybridisation; somatic mutation; gene therapy; PCR; primer; amplification; ss.

Synthetic

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                                                                                                                                                                                                                                                                                                                                                                                                                  This is the nucleotide sequence of a PCR primer used for amplification in suppressor (MTS) gene, to diagnose and treat cancer. The MTS gene is useful in the diagnosis and prognosis of human cancer, e.g. by standard nucleic hybridisation techniques, of patient samples. The mutated sequences are those that are present in somatic mutations of the gene in cancers. The vectors can be used for gene therapy strategies to replace construct protein in patients. These can also be used to the expression constructs also for therapeutic strategies. In addition the expression constructs can also be used to some the expression constructs can also be used to my. Recombinant MTS can be used to screen for drugs to be used for cancer therapy, and the protein itself may also be used to restore MTS
                                                                                                                                                                                                                                                                                                                 Nucleic acids based on multiple tumour suppressor, MTS, sequences - useful as hybridisation probes, primers and recombinant production of MTS in the diagnosis and treatment of cancers related to MTS mutation(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unmethylated CpG dinucleotide, immune response, bacterial meningitis, nacural, killer cell activation; NK cell; Th2 response; neonatal sepsis, pulmonary disorder; asthma; environmentally induced airway disease; bacterial infection; endocoxemia; therapy; cystic fibrosis; inflammatory bowel disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                     Example 12; Col 85-86; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unmethylated CpG dinucleotide 2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                505 GAGGGCTACCTGGAGAAGCT 524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 gaaggerrecresacaeser 1
                                                                                             94US-00214582,
94US-00215086,
94US-00215087,
94US-00227369,
94US-00251938,
95WO-US003316,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV47686 standard, DNA; 20 BP
                                                                95US-00480810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-US003678
                                                                                                                                                                                                                  (MYRI-) MYRIAD GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-NOV-1998 (first entry)
                                                                                                                                                                                                                                                                                    WPI; 1998-494842/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              function in a cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9837919-A1
                                                                07-JUN-1995;
                                                                                                                                                                 01-JUN-1994;
17-MAR-1995;
US5801236-A.
                                                                                                  18-MAR-1994;
                                                                                                                              18-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        J3-SEP-1998.
                                                                                                                                                  14-APR-1994
                                01-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV47686;
                                                                                                                                                                                                                                               Kamb A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1227
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This sequence represents an unmethylated CpG dinuclectide, and can be used in the method of the invention. The method is for treating a subject having, or at risk of having an acute decrement in air flow, comprising administering a nucleic acid sequence containing at least one administering a nucleic acid sequence containing at least one unmethylated CpG. The nucleic acids containing an unmethylated CpG this nucleic acids containing an unmethylated CpG thin the nucleic acids containing an unmethylated CpG thin cells (NK) or redirecting a subject s immune response from a Thic or Thi response by inducing monocytic and other cells to produce Thi cytokines. They can be used to treat pulmonary disorders having an immunologic component, such as asthma or environmentally induced airway positive bacterial infections or endotoxeemia including bacterial meningitis, neonatal sepsis, cystic fibrosis, inflammatory bowel disease and liver citrhosis, gram-negative abdominal abserces has mentalled to the preumonia, gram-negative abdominal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer; amplification; yeast; UAS; upstream activating sequence; UAS; transcription terminator; call cycle; Upstream Activation Sequence; UAS; promoter; phosphorylation; cyclin; cyclin-dependent kinase; CDK; vector; cyclin kinase inhibitor; CKI; growth; wound healing; cancer therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 abscess, haemorrhagic shock, disseminated intravascular coagulation, or an inflammatory response to lipopolysaccharide
                                                                                                                                                                       Use of nucleic acids containing an unmethylated CpG - for treating a subject having or at risk of having an acute decrement in air flow or inhibiting an inflammatory response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred, No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 0 A; 6 C; 14 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer #2 for human CDK2 codons 1-149.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             555 CCTCAGCCGCCCCTCCGTC 574
                                                                                                                                                                                                                                                     Claim 35; Page 27; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV60732 standard; DNA; 20 BP
                     28-FEB-1997; 97US-0039405P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-US018608
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96US-0031968P.
                                                      (IOWA ) UNIV IOWA RES FOUND.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
                                                                                               Schwartz DA, Krieg AM;
                                                                                                                                   WPI; 1998-480941/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BITT-) BITTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-1996;
27-NOV-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-APR-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bitter GA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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(ISIS-) ISIS PHARM INC.
  WPI; 1998-251302/22
                                                                                    Local Similarity
                                                                                                                                                      Key
modified_base
                                                                                                                                                                              14-APR-1998;
                                                                                                                         04-FEB-1999
                                                                                                                                                 Homo sapiens
                                                                                                                                                                   WO9846272-A1
                                                                                                                                                                                   14-APR-1997;
                                                                                      16;
                                                                                                                                              Synthetic
                                                                                                 20
                                                                                                                    AAV69958;
                                                                                 Query Match
                                                                                                         RESULT 1229
                                                                                    Best Loca
Matches
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AAV69949 to AAV69977 represent antisense oligonucleotides which are specifically hybridisable with a region of a nucleic acid encoding human For the antisense compound regalates the expression of the coligonucleotides which regulate the c-Jun protein. The present invention also describes antisense oligonucleotides which regulate the c-Jun protein. The antisense oligonucleotides are used for the diagnosis and treatment of diseases or disoraciers associated with Activating Protein 1 expression, of which c-Fos and c-Jun are subunits. The antisense oligonucleotides are used in compositions as c-Fos and/or c-Jun together with a carrier and a chemotherapeutic agent. They are used to regulate the expression of c-Fos or c-Jun in cells or tissues, preferably by inhibiting metastasis. They also regulate cell cycle expression and can be used to treat an animal
                                                                                                     Antisense oligonucleotides regulating Activating Protein 1 subunits - hybridise with c-fos and c-jun mRNA, used for regulating metastasis, cell cycle expression and hyperproliferative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA specific for Multiple Tumour Suppressor 1E1-beta gene - are useful for the diagnosis of cancers related to MTS1E1-beta mutation(s) and their treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MTS1; MTS1E1-beta; multiple tumour suppressor; diagnosis; cancer; germ-line mutation; familial melanoma locus; MLM; predisposition; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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80.0%; Pred. No. 9.3e+02;
rative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      with, or being prone to, a hyperproliferative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 0 A; 9 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human MTS1 and MTS1E1-beta PCR primer #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             725 AAGAGGGGCACCCTGCACC 744
                                                                                                                                                                                                                                                    Claim 5; Page 74; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 AAGGGGAGCAGCCGGCACC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV11263 standard; DNA; 20 BP
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94US-00215086.
94US-00215087.
94US-00227369.
94US-00251938.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 80.0 es 16; Conservative
                                  WPI; 1998-609906/51
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18-MAR-1994;
14-APR-1994;
01-JUN-1994;
17-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV11263;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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XX AAV1

XX AAV1

XX BE Huma

XX WEST Synt

XX WEST Synt

XX WEST SYNT

XX SYNT

XX SYNT

YX SYNT

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                                                                                                                                                                                                                                                                                                       brimers AAV60731-V60732 were used to PCR amplify codous 1-149 of the human cyclin-dependent kinase 2 (hCDX2) gene. The amplified product was used to generate a fusion protein comprising part of the hCDX2 sequence linked to codons 154.302 of the yeast PH085 gene. The fusion protein is used to codons 154.302 of the yeast PH085 gene. The fusion protein is used to screen for compounds that affect mammalian cell cycle regulatory proteins. The method comprises administering a compound to a cell line, which contains a reporter gene linked to an Upstream Activation Sequence (UAS) and a promoter, where the UAS binds a transcription control factor (UAS) and a promoter, where the UAS binds a transcription control factor (TCF) which is regulated through cyclin/cyclin-dependent kinase (CDK) phosphorylation. Also included in the construct is an effector gene providing a gene product to permit normal cyclin/CDK regulation of the TCF. Expression of the reporter gene is then analysed in the cell line, thereby determining whether the compound affects the normal regulation. The method can be used to identify inhibitors and activators of mammalian cell cycle regulatory proteins, especially inhibitors and activators of cyclin/CDK/CKI complexes. The identified agents can be used for stimulating growth of cells (as in wound healing), or regulating excessive cell growth and division (as in cancer therapy)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense oligonucleotide, phosphorothioate, regulation,
malignant tumour, cell cycle expression, hyperproliferative disease, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; c-fos; c-jun; activating protein 1; AP-1; diagnosis; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                 Screening for agents that effect cell cycle regulatory proteins - u cell line that expresses a reporter gene in response to regulation through phosphorylation by a cyclin/CDK system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
iive 0; Mismatches 4; Indels

    .20
    /*tag= a
    /note= "phosphorothioate linkages"

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 5 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Baker B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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                                                                                                                                                                                                                                         Example 4; Page 70; 93pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cacacriricascraeccaca 1
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Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
                                    Primers AAV11260-V11266 are used in the isolation of the human multiple tunmour suppression proteins, MTS1 and MTS1B1-beta. The MTS gene locus is also referred to as the familial melanoma (MLM) gene locus, located on human chromosome 9921. Germ line mutations in MTS genes can be used in the diagnosis of predisposition to cancers, e.g. melanoma, leukaemia, astrocytoma, gliboblastoma, lymphoma, glioma, Hodgkin's lymphoma, CLL, and cancers of the pancreas, breast, thyroid, ovary, uterus, testis, kidney, stomach and rectum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for detecting a selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.
                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                            Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
Example 12; Col 85-86; 72pp; English.
                                                                                                                                                                                                                                                                                                                                          505 GAGGCTACCTGGAGAGCT 524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 46; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                 20 GAAGGCTTCCTGGACACGCT 1
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98IL-00126627.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PTK 19 gene specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ18169 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-419113/35.
P-PSDB; AAY14704.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9934016-A2
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16-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ18169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vider B;
                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1231
                                                                                                                                                                                                                                                                                                                                                                                                                                                           $X88888888888888
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The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell; (c) determining in a selected gene family; and (c) calculating a performed and the methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for determining the cite of a selected treatment on a test cell. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired containing the pattern of gene expression in a selected genetic miling the pattern of gene expression in a selected gene family. Sequences AAZ17803-Z18342 represent primers that can be used
                                                                                                                                                                                                                                        ö
property. The method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AA217803-218342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes or cadherin superfamily genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.
                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                        ö
                                                                                                                                                                                             0.8%; Score 13.6; DB 1; Length 20;
30.0%; Pred. No. 9.3e+02;
ive 0; Mismatches 4; Indels
                                                                                                                                                        Sequence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                           1024 AAGCTGGCTGACTTTGGCCT 1043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 46; 102pp; English.
                                                                                                                                                                                                                                                                                                               AAGCTCGGGGACTTTGGGCT 20
                                                                                                                                                                                                                                                                                                                                                                                                                 B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-IL000625.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97IL-00122793
98IL-00126627
                                                                                                                                                                                                                  80.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PTK 18 gene specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                           AAZ18167 standard, DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                Local Similarity
les 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ18167;
                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vider B;
                                                                                                                                                                                                                                                                                                                                                                          RESULT 1232
                                                                                                                                                                                                                                      Matches
    88666666888
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Sequence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other;

SXS

superfamily genes or cadherin superfamily genes

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The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and the second cells. The method cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell; (c) determining a specified formula. The methods can be used for proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for determining the original cells capable of expressing an homeobox related dealired of containing cells capable of expressing an homeobox related dealired containing cells capable of expressing an homeobox related dealired geneting. Sequences AALT1803-Z18342 represent primers that can be used for the family. Sequences AALT1803-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genetic proximity, gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-FCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.
                                                                                                                                                                          Gaps
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                                                                                                                               0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
vative 0; Mismatches 4; Indel8
                                                                                               Seguence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                              1024 AAGCTGGCTGACTTTGGCCT 1043
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                                                                                                                                                                                                                                     AAZ18165 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PTK 17 gene specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                               Query Match
Best Local Similarity 80.09
Matches 16; Conservative
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P-PSDB; AAY14700.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
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                                                                                                                                                                                                                                                                                                                  RESULT 1233
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                                                                                                                                                                                                                                                                                     PAG; pregnancy associated glycoprotein; cattle; diagnosis; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New bovine polypeptides useful for early diagnosis of pregnancy
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Length 20;
                                                                                                                                                                                                                                                          Pregnancy associated glycoprotein (PAG) reverse primer B.
                              4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 1 A; 5 C; 4 G; 10 T; 0 U; 0 Other;
0.8%; Score 13.6; DB 1;
80.0%; Pred. No. 9.3e+02;
attive 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 52; 136pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   667 GGCAAAAGCAAAGCTCACAGA 686
                                                              1024 AAGCTGGCTGACTTTGGCCT 1043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 decahahahahahahah 1
                                                                                          1 AAGCTCGGGGACTTTGGGCT 20
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98US-0106188P.
                                                                                                                                                                    AAZ20188 standard; cDNA; 20
                                                                                                                                                                                                                                 05-JAN-2000 (first entry)
                   Local Similarity 80.0 nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Roberts RM, Green JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UMOR ) UNIV MISSOURI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-601132/51.
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28-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                19-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                     23-SEP-1999
                                                                                                                                                                                                                                                                                                                                                         Bos taurus.
                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                      AAZ20188;
        Query Match
                                                                                                                                            RESULT 1234
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RESULT 1235

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03-FEB-1999;
    12-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gamma-amino-butyric acid B receptor subunit; HG20; GABABR1a; depression;
epilepsy; neuropsychiatric disorder; dementia; muscular contraction;
central nervous system disorder; PCR primer; ss.
                                                                                               Human; raf; diagnosis; abnormal proliferative state; hyperproliferation;
cancer; psoriasis; blood vessel restenosis; c-raf kinase; antisense; ss.
                                                                                                                                                                                                                                                                                                                              Oligonucleotides targeted to human raf mRNA useful for treating and diagnosing abnormal proliferative states and inhibiting raf expression.
                                                                                                                                                                                                                                                                                                                                                                                  The invention provides antisense oligonucleotides targeted to mRNA encoding human raf and capable of inhibiting raf expression. The antisense oligonucleotides are useful for treating and diagnosing abnormal proliferative scates and hyperproliferation (e.g. cancer, psoritasis, or blood vessel restences), and inhibiting raf expression. Sequences AAZ11511-537 and AAZ11555-573 represent antisense oligonucleotides for human c-raf kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human GABA B receptor subunit HG20 PCR primer ngflt7-.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                        Human c-raf kinase antisense oligo ISIS # 5149.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1186 ATGGCCACAGGCCGTCCCCT 1205
                                                                                                                                                                                                                                                                                                                                                               Disclosure, Col 9, 29pp; English.
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            BP.
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95WO-US007111.
AAZ11521/c
ID AAZ11521 standard, DNA, 20
XX
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ID AAZ07001 standard; DNA; 20
                                                    (first entry)
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Best Local Similarity 80.0°
Matches 16; Conservative
                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                       Monia BP;
                                                                                                                                                                                                                                                                                                          WPI; 1999-527018/44.
                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                             31-MAY-1994;
31-MAY-1995;
                                                                                                                                                                                                          26-NOV-1996;
                                                    05-NOV-1999
                                                                                                                                                               USS952229-A
                                                                                                                                                                                     14-SEP-1999
                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                    Boggs RT,
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                                AAZ11521;
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Homo sapiens WO9940114-A1

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                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes two gamma-amino-butyric acid (GABA) B receptor (GABARR) subunits designated HG20 and GABARRIa. Cells expressing the new receptor abunits are useful for identifying GABARR agonists and antagonists. HG20 proteins and their antagonists are useful for inhibiting HG20 or GABARR function, useful for treating depression, eighlepsy, neuropsychiatric disorders. Generalism subscular contractions, and central nervous system disorders. The present sequence represents a PCR primer for human HG20, which is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, iPFK-2; cancer malignancy diagnostic assay; inflammatory disease;
inducible phosphofructokinase-2; tumour; malignant cancer; diagnosis;
therapy; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                Clark J;
                                                                                                                                                                                                                                                                          New DNA encoding human and murine receptor subunits, useful for identifying agonists and antagonists for treatment of depression, epilepsy and neuropsychiatric disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
vative 0; Mismatches 4; Indels
                                                                                                                                                                              Ng GYK, Kolakowski LF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                        (MERI ) MERCK & CO INC.
(MERI ) MERCK FROSIT CANADA INC.
(UTYTE-) UNIV TEXAS HEALTH SCI CENT SAN ANTONI.
(USSH ) US NAT INST OF HEALTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human iPFK-2 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                           Example 22; Page 85; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        901 ATGCACAAGGTGAAACTGTT 920
                                                                                                                                                                              Mcdonald T, Bonnert TP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PICO-) PICOWER INST MEDICAL RES.
(CHES/) CHESNEY J.A.
(MITC/) MITCHELL R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 AGGCAGCTGGAAACTGTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX58122 standard; DNA; 20 BP
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                                      98US-0073767P.
99WO-US002361.
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Best Local Similarity 80.0°
                                                                                                                                                                                                                                   WPI; 1999-527300/44.
                                      05-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-OCT-1998;
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                                                                                                                                                                            Liu Q, Mcd
Bonner II;
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invention relates to a cancer malignancy diagnostic assay for determining the presence of inducible phosphoffunctokinase-2 (ippr-2) specific sequences in a sample of a body or tumour fluid or tissue. The assay comprises obtaining a sample of a body or tumour fluid or tissue and performing a sequence identity assay to look for the presence of ippr-2 specific sequences. The method is useful for diagnosis of malignant orancer by detecting the presence of iPFK-2 specific sequences. Antisense iPFK-2 injouncebotides are useful for treatment of cancer and inflammatory disease. Antagonists of iPFK-2, such as an iPFK-2 enzyme inhibitor or anti-iPFK-2 antibody are also useful for treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; iPFK-2; cancer malignancy diagnostic assay; inflammatory disease; inducible phosphofructokinase-2; tumour; malignant cancer; diagnosis;
                                                                                                                                    This sequence represents a human iPFK-2 antisense oligonucleotide. The
                                                Cancer malignancy diagnostic assay useful for diagnosis of malignant cancer and, in treatment of cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cancer malignancy diagnostic assay useful for diagnosis of malignant cancer and, in treatment of cancer and inflammatory disease.
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                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human iPFK-2 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1679 CCAACTACATCTTCCCTGCT 1698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 4; Page 10; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PICO-) PICOWER INST MEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 ccaacecarcrrcececr 20
                                                                                                  Claim 4; Page 13; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX58144 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97US-00961578.
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                                                                                                                                                                                                                                                                                                                           and inflammatory disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MITC/) MITCHELL R A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CHESNEY J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-313301/26.
              WPI; 1999-313301/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX58144;
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This sequence represents a human iPFK-2 antisense oligonucleotide. The invention relates to a cancer malignancy diagnostic assay for determining the presence of inducible phosphofructokinase-2 (iPFK-2) specific

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Gaps

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sequences in a sample of a body or tumour fluid or tissue. The assay comprises obtaining a sample of a body or tumour fluid or tissue and performing a sequence identity assay to look for the presence of iPFK-2 specific sequences. The method is useful for diagnosis of malignant cancer by detecting the presence of iPFK-2 specific sequences. Antisense inflammatory disease. Antagonists of iPFK-2, such as an iPFK-2 enzyme inflammatory disease. Antagonists of iPFK-2, such as an iPFK-2 enzyme and inflammatory disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV74237-V74253 are oligodeoxynucleotide (ODN) primers used to describe a method for enhancing the immunostimulatory effect of an antigen encoded involves determining the cpd-N and Cpd-S motife present in the construct. The method involves determining the Cpd-N and Cpd-S motife present in the construct, removing neutralising CpG (Cpd-N) motife and optionally inserting attimulatory CpG (Cpd-S) motife in the construct, thereby producing a nucleic acid construct having enhanced immunostimulatory efficacy. The method can be used for immunisation against viral antigens, e.g. from hepatitis B virus (HBV), bacterial antigens or an antigen derived from a parasite. They can also be used for expression of a therapeutic polypeptide, e.g. growth factors, toxins, tumour suppressors, cytokines,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CpG-N motif; immunostimulation; antigen; CpG-S motif; immunisation; ODN; viral antigen; bacterial antigen; parasite; therapeutic; growth factor; toxin; tumour suppressor; cytokine; apoptotic protein; interferon; hormone; clotting factor; liqand; receptor; oligodeoxynucleotide; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            - for
the
                                                                                                                                                                                                                                                                            0; Gaps
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                                                                                                                                                                                                                                  0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
ative 0; Mismatches 4; Indels
                                                                                                                                                                                                Sequence 20 BP; 4 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wu T;
                                                                                                                                                                                                                                                                                                                  1679 CCAACTACATCTTCCCTGCT 1698
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-US010408.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV74243 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CpG-N motif O-ODN 2001 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                       Local Similarity 80.0 nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Davis HL, Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (QIAG-) QIAGEN GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-059712/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-1998;
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15-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                        Query Match
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AAV74294/

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Oligonuclectides AAV70607-11 were used to isolate nucleic acid encoding a murine multiple tumour suppressor IEI-beta (WTSIEI-beta) protein. Primers designed from the gene can be used to design primers to detect abnormalities i.e. polymorphisms which may predispose towards malignancies such as melanoma, leukaemia, astrocytoma, lymphoma, glioma, as well as tumours of e.g. the breast, thyroid, pancreas, uterus and kidneys. (Updated on 20-MAR-2003 to correct PR field.) (Updated on 20-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse multiple tumour suppressor gene segment - useful for primer design.
                                                                             Human; multiple tumour suppressor 1 gene; MTS1; cancer; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4; Indels
                                               PCR primer used to isolate murine MSTIE1-beta gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.6; DB 1;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 13; Col 53; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 505 GAGGGCTACCTGGAGAAGCT 524
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                                                                                                                                                                                                                                                                              95WO-US003316.
                                                                                                                                                                                                                                                                                                                                   (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ02575 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                     Jiang P, Kamb A, Stone S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-0CT-1999 (first entry)
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Best Local Similarity 80.0°
                 (first entry)
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Chlamydia trachomatis.
 (revised)
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20-MAR-2003
03-FEB-1999
                                                                                                                                                                                                                                            28-JUL-1995;
                                                                                                                                                                                                                                                                              17-MAR-1995;
07-JUN-1995;
                                                                                                                                                                          US5843756-A.
                                                                                                                                                                                                              01-DEC-1998
                                                                                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                            ICAM-1; intercellular adhesion molecule-1; antisense; primer; prevention; perfusion injury; transplantation; pre-operative treatment; donor; organ;
apoptotic proteins, interferons, hormones, clotting factors, ligands and receptors. (Updated on 20-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of antisense oligonucleotide against ICAM-1 - for preventing perfusion injury during transplantation of e.g. kidney, heart, lung or
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ilarity 80.0%; Pred. No. 9.3e+02;
Conservative 0; Mismarch...
                                                                                     0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels
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                                                      Seguence 20 BP; 0 A; 6 C; 14 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                              ICAM-1 antisense oligonucleotide primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                               555 CCTCAGCGGCGCCTCCGTC 574
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97DE-01045666
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                                                                                                                            16; Conservative
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Matches 16; Conserv
                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pancreas.
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                                                                                          Query Match
Best Local
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Length 20;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                        Genome sequence of Chlamydia trachomatis.
                                                                                                                                                                                                       Disclosure; Page 1536; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1387 CTCCTCACCAAGCTGTTGCA 1406
 97FR-00015041.
97FR-00016034.
98US-0107077P.
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                                                                                                                                       WPI; 1999-371125/31,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                    (GEST ) GENSET
28-NOV-1997;
17-DEC-1997;
04-NOV-1998;
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                                                                                                   Griffais R;
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Matches
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PCR primer used to amplify an ORF of Chlamydia trachomatis.
CTCCGAACAAGCTGTTCCA 20
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                                                                                                                                                                                               RESULT 1243
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Vaccine, eye disease, conventional trachoma, nonendemic trachoma, paratrachoma, inclusion conjunctivitis, genital disease, perihepatitis, nongonococcal uretritis, epidymitis, cervicitis, salpingitis, PCR primer, bartholinitis; pneumopathy, venereal lymphogranulomatosis, ss. Chlamydia trachomatis

WO9928475-A2

10-JUN-1999,

98WO-IB001939

27-NOV-1998;

97FR-00015041. 97FR-00016034. 98US-0107077P. 28-NOV-1997; 17-DEC-1997; 04-NOV-1998;

GEST) GENSET

Griffais R;

WPI; 1999-371125/31

Genome sequence of Chlamydia trachomatis.

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PCR primers AAZ01426-Z06209 were used to amplify open reading frames choose polypeptides (see AAY36754-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachom, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, practicis, genital diseases such as nongonococcal uretritis, prehompathy in breast feeding infants; and venereal lymphogramulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                  Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                      Disclosure; Page 1447; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                               0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 80.0
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CGCG 106

87 CGGCTCTGAGGTTGC

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Vaccine, eye disease, conventional trachoma, nonendemic trachoma, paratrachoma, inclusion conjunctivitis, genital disease, perihepatitis, nongonococcal utetritis, epidymitis, cervicitis, salpingitis, PCR primer; bartholinitis, pneumopathy, venereal lymphogramulomatosis, ss. PCR primer used to amplify an ORF of Chlamydia trachomatis. BP. AAZ05818 standard; DNA; 20 (first entry) 07-0CT-1999 AAZ05818;

97FR-00015041. 97FR-00016034. 98US-0107077P. 98WO-IB001939 Chlamydia trachomatis. (GEST) GENSET 27-NOV-1998; 28-NOV-1997; WO9928475-A2 17-DEC-1997; 04-NOV-1998; 10-JUN-1999 Griffais R; Synthetic.

Genome sequence of Chlamydia trachomatis. Disclosure; Page 1802; 1755pp; English

WPI; 1999-371125/31.

PCR primers AAZ01426-206209 were used to amplify open reading frames (OREs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY36754.737949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, eg. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs accode polygeptides (see AAX36754-737949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal urstrilis, epidymitis, cervicitis, salpingitis, perimpgatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis.
                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccine, eye disease, conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease, perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                 pneumopathy in breast feeding infants; and venereal lymphogranulomatosis.
The polypeptides of the invention may be of use in treating these
                                                                                                                                              Gaps
cervicitis, salpingitis, perihepatitis, bartholinitis;
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0
                                                                                                          0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
ative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                      PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                                           Sequence 20 BP; 2 A; 12 C; 1 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 1536; 1755pp; English.
                                                                                                                                                                             1636 AGGCAGCGGCTGGAGGGATG 1655
                                                                                                                                                                                                            20 AGGGAGAGGCGGGAGTGTG 1
                                                                                                                                                                                                                                                                                          AAZ02583 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chlamydia trachomatis.
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                                                                                                                             Local Similarity
ses 16; Conserv
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17-DEC-1997;
04-NOV-1998;
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   epidymitis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
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This oligonuclectide represents an antisense oligonuclectides (ASO) targeted to a region in the intercellular adhesion molecule (ICAM)-1 genewhich is generated by a method of selectively ranking mucleic acid molecules for inhibitory efficiency. The method comprises: (a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA.ASO-DNA hybrid nucleic acid base pair types in a target sequence zone RNA.ASO-DNA hybrid nucleic fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions, and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Selectively ranking nucleic acid molecules, for inhibitory efficiency comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Target; antisense; selective rank; inhibition; ranking; stability; interaction; intercellular adhesion molecule; ICAM; ss.
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/trag= a
/note= "contains phosphorothioate internucleotide
linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide ISIS#1939 targeted to ICAM-1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         226 GAGAGTGGTGGTGGCGG 245
283 GGGGAACTTCGTTCTGCACG 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 chchcccchactccrccccc 1
                                                             20 decentricerrierre
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Matches 16; Conservative
                                                                                                                                                                                                                                       standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-105098/09.
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                                                                                                                                                                                                                                                                                                                                                                         30-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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XX AAX 06531/c

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Gaps

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Cuery Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels

Sequence 20 BP; 9 A; 7 C; 3 G; 1 T; 0 U; 0 Other;

Synthetic.

AAX21345;

RESULT 1247 AAX21345

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CER primers AAZ10708-33 were used to amplify exons I to 11 of human HKNGI

(Hong Kong new gene 1). HKNGI is a gene associated with bipolar affective
disorder (BAD). HKGNI polymucleotides are useful to identify compounds

modulating HKNGI gene expression or HKNGI polymeptide expression/
activity. Compounds inhibiting or enhancing HKNGI gene expression or
activity. Compounds inhibiting or enhancing HKNGI gene expression or
activity in individuals can then be administered therapeuticially to treat
HKNGI-mediated disorders, especially neuropsychiatric disorders e.g. BAD,
schizophrenia, or HKNGI-mediated myopia disorders, such as early-onset
contropic of HKNGI-mediated myopia disorders is uch as early-onset
contropic to identify individuals having, or at risk of developing, HKNGI
chagnosis to identify individuals having, or at risk of developing, HKNGI
confediated disorders due to mutations in the HKNGI gene. Such mutations
confediated by result in the production of a protein with a different
sequence to the human full-length HKNGI polypeptide or splice variant
sequence to the human full-length HKNGI polypeptide or splice variant
sequence to the human full-length HKNGI polypeptide or splice variant
sequences, especially the substitution of a lysine for a glutemic acid at
confediated special substitution of a lysine for a glutemic acid at
to produce probes or primers to identify similar sequences (e.g. mutants
confediated sequences from different species) and to produce transgenic animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; alpha-7 nicotinic receptor; neuronal; hybridisation; probe;
alpha-7 neuronal nicotinic acetylcholine receptor; schizophrenia;
HKNG1; Hong Kong new gene 1; bipolar affective disorder; BAD; neuropsychiatric disorder; early-onset autosomal dominant myopia; schizophrenia; splice variant; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New HKNG1 polynucleotides useful in diagnosis and treatment of neuropsychiatric disorders, e.g. bipolar affective disorders and
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98US-0088312P.
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Guery Match
Bust Local Similarity 80.v.,
Bust Local 16; Conservative
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(REGC ) UNIV CALIFORNIA.
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                                                                                                                                  Homo sapiens.
                                                                                                                                                                                W09947535-A1
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22-JAN-1999;
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                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX56166,
            8x566666666666666668x8x34445x8xx356666666666666666
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primers AAX21344-X21345 were used to PCR amplify a 375 bp fragment of the apolipoprotein E gene. The invention relates to the diagnosis of Alzheimer's disease (AD) by detecting one or murations, in the genomic region that regulates expression of the apolipoprotein E (apo E) that results in: (a) altered sene expression relative to a control population, or (b) altered relative expression relative to a control Alternatively AD is detected by determining the levels of apo E. Alternatively AD is detected by determining the levels of epsilon-2, -3 or -4 alleles or mutations in the Thi/E470s sequence (a polymorphism in the promotter region). The T allele of Thi/E470s is associated with increased risk of AD (independently of the effect of the epsilon allele) and increases the risk associated with epsilon 4 in epsilon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Diagnosing Alzheimer's disease - by detecting mutations in the regulatory region of the apo E gene, or levels of apo E allele expression.
                                                                                                                                                                                                                                                                                              Primer; PCR; amplification; apolipoprotein B; human; brain; diagnosis;
Altheimer's disease; mutation; gene expression; polymorphism; promoter;
allele; heterozygote; 8s.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
.ive 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                             Primer #2 for amplifying apolipoprotein E gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (INSP ) INST PASTEUR LILLE.
(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chartier-Harlin M, Lambert J, Amouyel P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 699 ACTCAAGGAGATCAGACTGG 718
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                                                                                                                   AAX21345 standard; DNA; 20
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                                                                                                                                                                                                                 (first entry)
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ses 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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Gaps

AAZ10728;

XEXEXEX

RESULT 1248

AAZ10728/

Query Match

ଚ 셤 schultz621-3.rng

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The present invention describes an isolated nucleotide sequence (I) encoding at least a portion of the human alpha-7 neuronal nicotinic acceptor (alpha7-hnAchR). Also described are: (I) a peptide encoded by (I); (2) a vector comprising (I); (3) a host cell transformed with a vector of (2); (4) a polymucleotide comprising at least 15 nucleotides which hybridises under stringent conditions to at least a portion of (I); (3) a method for detection of a polymucleotide encoding alpha 7-hnAchR in a biological sample; and (6) a method for amplification of an of nucleic acid from a sample suspected of containing nucleic acid contains and probes from the present encoding alpha 7-hnAchR. The primers and probes from the present condition can be used on brain tissue and blood samples of humans suspected of suffering from schizophrenia, small cell lung carcinoma, breast cancer and nicotine-dependent illness. This is particularly useful condidiagnosed are epilepsy (e.g. juvenile myoclonic epilepsy) and prader-Willi and Angelman's syndromes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumour necrosis factor alpha; TNF-alpha; antisense oligonucleotide; ASO; inhibition; expresssion; treatment; disease; disorder; ss.
small cell lung carcinoma; breast cancer; nicotine-dependent illness;
epilepsy; juvenile myoclonic epilepsy; Prader-Willi syndrome;
Angelman's syndrome; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                  Human alpha-7 neuronal nicotinic acetylcholine receptor and related polynucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 13.6; DB 1; Length 20;
llarity 80.0%; Pred. No. 9.3e+02;
Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tumour necrosis factor alpha antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 8 A; 9 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; Page 63; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    218 GCCTGGATGAGAGTGGTGGT 237
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                                                                                                                                                                                                                                                                                                      Leonard S, Freedman R;
                                                                                                                                                                                                                                                                                                                                          WPI; 1999-288306/24.
                                                                                                                                                                                                                                                    (LEON/) LEONARD S. (FREE/) FREEDMAN R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
nes 16; Conserv
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                                                                                                                                                                                    15-OCT-1998;
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                                                                                     Homo sapiens
                                                                                                                    WO9920757-A2
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                                                                     Synthetic.
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Antisense oligonuclectides (ASO) for inhibiting a tumour necrosis factoralpha (TNP-alpha) gene in an animal, preferably a human, comprise 12-50 nucleotides, 90% of which are complementary to a region of mENA containing a GGGA sequence motif. The ASO is used to inhibit expression of a gene in an animal and for treating the animal when afflicted with a fisaeae or disorder characterised by the presence of an mRNA from a gene containing a GGGA motif. The ASO are specifically targeted to a GGGA sequence motif found in mRNA from a gene. A study of known ASO has shown that at least half of the most effications ASO's contain one or more TCC motifs. This ASO comprises a TCCC motif followed by a cytosine residue and corresponds to a region of the human ICAM-1 3' untranslated region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                                                                                                                                                                                             Generation of antisense oligonucleotides - by specifically targeting a GGGA motif found in mRNA sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                               Example 2; Page 37; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 dadadegeaagreeregees 1
                                                                                              (UYJE-) UNIV JEFFERSON THOMAS
                                98WO-US013711.
                                                                97US-0051705P.
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                                                                                                                                                              WPI; 1999-105767/09.
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                                                                                                                                Tu G, Israel Y;
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                                02-JUL-1998;
                                                                03-JUL-1997;
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14-JAN-1999
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Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;

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AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX34564-AAX35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae mucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae
                                            AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae present of Chlamydia pneumoniae present of Chlamydia pneumoniae present of Chlamydia pneumoniae present of sease such as pereumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema rodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX14584-AAX15879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae especially where the vector directs the expression of a neutralising epitope of C. pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                                                                                                                                                                                                                                                                                                               0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
.ive 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 4 C; 6 G; 5 T; 0 U; 0 Other;
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                  Page 1787; Disclosure; 1912pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                953 GCCACCGCCAGAAGGTGCTA 972
                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GCTATCGGCAGATGATGCTA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX92771 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 80.0'
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chlamydophila pneumoniae
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AAX92771/c
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                                                                                                                                                                                                                                                                                     Respiratory disease, pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
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80.0%; Pred. No. 9.38+02;
tive 0; Mismatches 4; Indels
Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indel8
                                                                                                                                                                                                                                                             PCR primer used to amplify an ORF of Chlamydia pneumoniae.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genome sequence of Chlamydia pneumoniae.
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                                                                405 GTCTCCAGTGAGAGTGCGTA 424
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                                                                                                                                                                       AAX94323 standard; DNA; 20 BP.
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   Query Match
Best Local Similarity 80.0%;
Matches 16; Conservative
                                                                                                                                                                                                                                   (first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                     AAX94323;
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                                                                                                                                           RESULT 12
AAX94323/
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Gaps ô RESULT 1254 AAX94068/c

AAX94068

Synthetic

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AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (See AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent citis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX34584- AAX35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae especially where the vector directs the expression of a neutralising epitope of C. pneumoniae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                     Genome sequence of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                         Page 1849; Disclosure; 1912pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                761 CCCTGCTCAAGGACCTCAAA 780
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AAX96621 standard; DNA; 20
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Matches 16; Conserv
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                                                                                                                                                                                      GEST ) GENSET
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04-NOV-1998;
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                                                     03-JUN-1999
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sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine;
neutralising epitope; PCR primer; ss.
                                                                                                                                             Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
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                                                                                                              PCR primer used to amplify an ORF of Chlamydia pneumoniae.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genome sequence of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 1641; Disclosure; 1912pp; English.
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  AAX94068 standard; DNA; 20
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Chlamydophila pneumoniae.
                                                                           (first entry)
                                                                                                                                                                                                                                                   Chlamydophila pneumoniae.
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epitope of C. pneumoniae
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les 16; Conserv
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04-NOV-1998;
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AAX96741;

RESULT 1255

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Query Match

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AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX34584- AAX35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae especially where the vector directs the expression of a neutralising
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                                                                                                                                                                                                                                                                                                                                                                                          4; Indels
                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 5 A; 2 C; 8 G; 5 T; 0 U; 0 Other;
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                              Genome sequence of Chlamydia pneumoniae.
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                                                                   Page 1840; Disclosure; 1912pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                               542 TCTTTGACAAGCCCCTCAGC 561
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98US-0107078P.
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les 16; Conservative
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                                                                                                                                                                                                                                                                                  epitope of C. pneumoniae
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WPI; 1999-357842/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA08849-60 are oligonucleotide primers and adaptors used in cloning the murine DKR-3 gene (AAA08838). The murine DKR-3 open reading frame has bomology to human rig-11ke 7-1 murine DKR-3 open reading frame has elest 4 gene. DKR-1 is a human ortholog of dkk-1 (dickkopf-1), a novel gene identified in Kenopus and mouse, purportedly an arragonist of wnt-8 pattern. DKR-2, -3 and -4 are each related to DKR-1 by their cysteine pattern. DKR-1; a lao involved in morphogenesis in the developing embryo, and therefore a growth factor. By inference DKR polypeptides are also growth factors. The DKR polypeptides are useful for treating cancer, e.g. mammary tumors, stem cell tumors, or other cancers in which the wnt and/or soulc hedgehog (shh) signal transduction pathways are activated. They can also be used to enhance tissue differentiation, such as bone formation and hematopoietic cell formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DKR-3; human rig-like 7-1 mRNA; chicken lens fiber protein; clfest 4; dkk-1; dickkopf-1; antagonist; wnt-8 signaling; morphogenesis; primer; growth factor; cytostatic; sonic hedgehog; tissue differentiation; ss.
nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae {\bf p}
                                                                                                                                                                 Gaps
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                                                                                                                   V Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Fred. No. 9.3e+02; Pes 16; Conservative 0; Mismatches 4; Indels
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    RACE nested primer for murine DKR-3 cDNA synthesis.

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                                                                                Sequence 20 BP; 6 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                      9 GCGTAAAGGATGGACAGGAA 28
                                                                                                                                                                                                                                              1 GCGTTCAGGATCTACAGGAA 20
                                                                                                                                                                                                                                                                                                                                              AAA08858 standard; DNA; 20 BP
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                                                                                                                          Query Match
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1 AACATGCAGCGGCTCGGGGG 20

10-APR-2000 (first entry)

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The invention relates to the isolation of the gene encoding the human multiple tumour suppressor 1 (MTS1) (AAA95633). The MTS1 protein has a cytostatic activity and is used in protein replacement therapy. This sequence is a PCR primer used to isolate the mouse pl6 gene (a homologue of the MTS1 gene). MTS1 is useful in diagnosing human cancers such as (ocular) melanoma, leukemia, astrocytoma, glioblastoma, lymphoma, glioma, Hodgkin's lymphoma, multiple myeloma, sarcoma, myosarcoma, cholangiocarcinoma, squamous cell carcinoma, CLL, and cancers of pancreas, breast, stomach, brain, proteter, bladder, thyroid, ovary, uterus, testis, kidney, colon and rectum. The MTS1 gene and protein is useful in gene therapy, protein replacement therapy and protein mimetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel protein composition useful in protein replacement therapy for diagnosing and treating cancer comprises a specific weight percent of human multiple tumor suppressor I polypeptide.
                                                                                                                                                             Cytostatic; human; multiple tumour suppressor 2; MTS2; diagnostic; cancer; gene therapy; protein replacement therapy; ss; mouse.
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94US-00215087.
94US-00227369.
94US-00251938.
                                  AAA95656 standard; DNA; 20 BP
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95US-00480810.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MYRI-) MYRIAD GENETICS INC.
                                                                                               14-FEB-2001 (first entry)
                                                                                                                                Mouse P16 gene primer #1.
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07-JUN-1995;
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14-APR-1994;
                                                                                                                                                                                                                                                                         18-JUL-2000.
                                                                                                                                                                                                                                                                                                                                       18-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                     01-JUN-1994
                                                                 AAA95656;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ57446;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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AAZ57446/c
ID AAZ57446
XX
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Kamb A;
                                                                                                                                                                                                              Mus sp.
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The present invention describes oligonuclectides containing bioreversible phosphate ester groups and their mimetics. The oligonuclectides are of value in therapeutics, diagnostics, and as research reagence. The compound from the present invention may be used in control of hereditary, metabolic, and/or cellular processes in any organism utilising DNA-RNA cranscription and/or RNA-protein translation. These organisms include prokaryotic undelular and multicellular organisms; including bacteria, yeasts, protozoa, algae, and all plants and higher animal forms, including warm blooded animals, particularly unans, also corganelle sub-cellular translation and transcription processes. The new synthetic process provides pro-oligonucleotides, i.e., oligonucleotides blocked at phosphate groups by bioreversible groups which can be cleaved by intracellular and intercellular and drugs. By careful selection of protecting groups, deprotection of mucleobases and partial deprotection of protecting groups, deprotection of mucleobases and partial deprotection of specific groups include s-pivaloylmercaphocethy! (SBMS) and complete sub-oligonal process and partial deprotection cyanocthylcarbony! (CEOC) groups. Spacer molecules include diglycoly!

CCCCH_ZOCH_ZOCH_ZOC) and its analogue with a catechol biresidue replacing the cyanom and its analogue with a catechol biresidue replacing the cyanom timent invaries and biosphoteto acid). The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                 Phosphorothioate; antisense oligonucleotide; triester oligonucleotide; bioreversible phosphate blocking group; therapeutic; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide bioreversible phosphate esters used as, e.g. research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred, No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                              /*tag= a
/note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                           Phosphorothicate oligonuclectide SEQ ID NO:5.
                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Page 35; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 dácadadadadrogradada 1
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                                                                                                                                                                                                                                                                                                                                                                                    99WO-US013141.
                                                                                                                                                                                                                                                                                                                                                                                                                            98US-00095822.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Guzaev A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-116518/10.
                                                                                                                                                                                                    Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                  10-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                            11-JUN-1998;
                                                                                                                                                                                                                                                                                                     WO9964434-A1
                                                                                                                                                                                                                                                                                                                                             16-DEC-1999.
                                                                                                                                                            Synthetic.
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Peroxisome proliferators are agents that induce peroxisomal proliferation. Peroxisome proliferator-activated receptors (PPARS) are members of the steroid receptor superfamily, and are ligand-activated transcription factors. There are three mammalian subtypes of PPAR, alpha, beta (also known as delta) and gamma. The present invention relates to inhibiting angiogenesis by contacting PPAR gamma with a PPAR gamma characterised by excessive neovascularisation e.g. rheumatoid arthritis, psoriasis, atheroselerosis, diabetes, retinopathy, retrolental fibroplasia, neovascular glaucoma, age-related macular degeneration, transplantation, chronic inflammation, und inflammation, endometriosis and obesity. The present sequence is a PCR primer for human PPARbeta gene. The resulting PCR product was used in the analysis of PPARbeta
                                                                                                         neovascularisation; tumour growth; metastasis; rheumatoid arthritis; psoriasis; atherosclerosis; diabetes; retinopathy; PPAR; Human; retrolental fibroplasis; age-related macular degeneration; neovascular glaucoma; hemanda, thyroid hyperplasis; Grave's disease; inflammatiobesity; transplantation; transcription factor; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Endocrine disruptor; dioxins; organic halocarbon; phenol; agrochemical; phthalate esters; aromatic hydrocarbon; organotin compound; oestrogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Inhibiting angiogenesis comprises administration of PPAR gamma ligands for treating tumors, neovascularization and rheumatoid arthritis.
                                                                                       Peroxisome proliferator activated receptor; angiogenesis inhibition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
rative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer for p38 nucleotide sequence amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 1 A; 6 C; 4 G; 9 T; 0 U; 0 Other;
                                                    Human PPARbeta gene reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 31; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    128 ATCGGATGAAGAAGATCAAA 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 AGCGGATCAAGAAGACCGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA59793 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                 99WO-US027612.
                                                                                                                                                                                                                                                                                                                                                                    98US-01109328P.
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                   (first entry)
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les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gerritsen ME, Xin XE;
                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-411807/35.
                                                                                                                                                                                                                                                         WO200030628-A2.
                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                 18-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                  20-NOV-1998;
20-JAN-1999;
                   17-OCT-2000
                                                                                                                                                                                                                                                                                            32-JUN-2000.
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Matches
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24445444
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disruptor is new and comprises isolation of mRNA from cells, tissue or organized must be endocrine disruptor, and hybridising with a DNA array contact with the endocrine disruptor, and hybridising it with a DNA array containing immobilized gene fragments from genes which may be affected by the endocrine disruptor. The results of the hybridisation are then compared with a compariation sample to of the hybridisation are then compared with a compariation sample to detect genes whose expression is altered by endocrine disruptors such as altered panes whose expression is altered by endocrine disruptors such as disruptor incompounds, ocstrogens, mylex, toxaphene, aldicarb and kepones. The types of genes whose expression may toxaphene, aldicarb and kepones. The types of genes whose expression may be altered by these disruptors include those involved in nuclear receptor transcriptional coupling, kinase type signal transduction, gonad differentiation, receptor type kinases, intermediate filament markers, apportosis, DNA damage response, repair and recombination, receptors, cell cycle and growth regulators, cell adhesion, motility and invasion, angiogenesis regulation, invasion regulation, cell-cell interaction, RNO family small GTPase regulation and growth factors and cytchines.

Completed sequences of genes which may be affected by an endocrine
            nuclear receptor transcriptional coupling; gonad differentiation, intermediate filament marker; cell cycle; growth; regulation; oncogene; tumour suppressor; apoptosis; DNA damage response; cell adhesion; motility; angiogenesis regulation; invasion regulation; growth factor; cytokine; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                             mRNA from calls exposed to an endocrine disruptor is hybridized with a DNA array of gene fragments for detection of genes whose expression is altered by the endocrine disruptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
mylex; toxaphene; aldicarb; kepones; kinase signal transduction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                        Kimizuka F, Kato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 5 A; 5 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer for mouse beta coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 63; 81pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1236 ACACTICATCTTCCGTATCT 1255
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                                                                                                                                                                                                                                                                                                                                                                        Mineno J,
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                                                                                                                                                                                                                                                       99WO-JP005964.
                                                                                                                                                                                                                                                                                           98JP-00310285.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   80.08;
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                                                                                                                                                                                                                                                                                                                                (TAKI ) TAKARA SHUZO CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                        Kondo A, Sagawa H,
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-365642/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                          WO200026404-A1.
                                                                                                                                                                                                                                                       28-OCT-1999;
                                                                                                                                                                                                                                                                                           30-0CT-1998;
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                                                                                                                                                                                                                11-MAY-2000
                                                                                                                                     Synthetic.
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This sequence is a PCR primer for DNA encoding mouse beta, protein which is homologous to the human MYSIBlbera protein. The invention relates to a method for diagnosing a polymorphism associated with a predisposition to cancer by detecting a germ-line alteration of a wild-type Multiple Tumour Suppressor (MYS) gene or its expression products in a human sample. The method comprises detecting a germ-line alteration of a wild-type MXS gene or its expression products in a human sample, the alteration indicating a predisposition to at least one of the cancers. The cancer is selected from melanoma, leukaemia, astrocytoma, glioblastoma, lymphoma, glioma, Hodgkin's lymphoma, chronic lymphocytic leukaemia (CLL), and cancers of the pancreas. Dreast, thyroid, ovary, uterus, testis, kidhey, stomach and rectum. The method may be used as the basis for developing very important diagnostic tests capable of predicting the predisposition to cancer. The MYS gene is involved in the progression of multiple tumour types and may provide means for a general anti-cancer therapy by virtue of its ability to suppress tumour growth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Multiple tumour suppressor; MTS2; human; diagnosis; Hodgkin's lymphoma; cancer predisposition; melanoma; leukaemia; lymphoma; glioma; MTS1Elbeta; PCR primer; ss.
MTS; human; polymorphism detection; cancer predisposition; astrocytoma; Multiple Thumour Suppressor gene; melanoma; leukaemia; glioblastoma; lymphoma; glioma; Hoddkin's lymphoma; chronic lymphocytic leukaemia; therapy; MTSIElbeta; PCR primer; sa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Diagnosing a polymorphism associated with a predisposition for cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer for human MTS1Elbeta 1 coding sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                  Skolnick MH, Cannon-Albright LA, Kamb A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 12; Col 50; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            505 GAGGGCTACCTGGAGAAGCT 524
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94US-00215086.
94US-00215087.
94US-00227369.
94US-00251938.
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                                                                                                                                                                                                                                                                                                                                           95US-00474083.
                                                                                                                                                                                                97US-00848251.
                                                                                                                                                                                                                                                                                                                                                                              (UTAH ) UNIV UTAH RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                               (MYRI-) MYRIAD GENETICS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-070785/06
                                                                                                                                                                                                29-APR-1997;
                                                                                        Homo sapiens
                                                                                                                           US5989815-A.
                                                                                                                                                                                                                                                                    8-MAR-1994;
                                                                                                                                                                                                                                                                                                                                           07-JUN-1995;
                                                                                                                                                                                                                                                                                       14-APR-1994;
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17-MAR-1995
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                                                                                                                                                              23-NOV-1999
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This sequence represents a PCR primer for the human multiple tumour suppressor IEIbeta (MTSIEIbeta) coding sequence. The invention relates to the human MTS2 DNA and protein sequences. The DNA sequences are useful to diagnosing or determining a predisposition to cancers e.g. melanoma, leukaemia, lymphoma, glioma, Hoddxin's lymphoma and cancers of the pancreas, breast, thyroid, ovary, kidney, uterus and stomach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DBL gene; Duffy-binding like gene; ebl-1; Duffy Antigen Binding Protein; DABP; Sialic Acid Binding Protein; SABP; malaria; vaccine; immunisation; protozoacide; eba-175; PCR primer; 88.
                                                                                                                                                                                                                                                                                            Multiple tumor suppressor cDNA, useful for diagnosing or determining a predisposition to cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Miller LH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Plasmodium DBL family conserved motif isolating primer UNIEBPSA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.38+02;
iive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wellems TE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peterson DS, Su X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                 Example 12; Col 50; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            505 GAGGCTACCTGGAGAGCT 524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 GAAGGCTTCCTGGACACGCT 1
                                                                                                                                       94US-00215086.
94US-00215087.
94US-00227369.
94US-00251938.
95WO-US003316.
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                                                                                                  95US-00486047
                                                                                                                          94US-00214582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95US-00487826
                                                                                                                                                                                                                  (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match
Best Local Similarity 80.09
Matches 16; Conservative
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                                                                                                                                                                                                                                                                   WPI; 2000-038259/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmodium sp.
                       Homo sapiens.
                                                                                                                                                              14-APR-1994;
01-JUN-1994;
17-MAR-1995;
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                                                                                                                         18-MAR-1994;
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                                               US5994095-A.
                                                                        30-NOV-1999,
           Synthetic
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                                                                                                                                                                                                                                           Kamb A;
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(ISIS-) ISIS PHARM INC.
                                                                                                                                            Mehta R, Hardee GE,
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                                                                                                                     WO9960167-A1
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                                                                                                                                   21-MAY-1998;
                                                                                         07-MAR-2000
                                                                                                                         25-NOV-1999.
                                                                                                                Synthetic.
                                                                                     AAZ48638;
                                                                           RESULT 12
AAZ48638/
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WPI; 2000-349676/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-JUL-1998;
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08-DEC-1997;
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18-MAR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6060301-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-MAY-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA39378;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kamb A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA39378,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8588888888888888888888
                                                                                                                                                                                                                                                                                                                                                                                                                    8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense inhibitor; oligonucleotide delivery agent; erythema multiforme; expression modulator; cellular adhesion protein; malignant melanoma; cellular proliferation modification; toxic epidermal necrolysis; psoriasis; lichen planus; carcinoma; Paget's disease; Kaposi's sarcoma; pulmonary fibrosis; Lyme disease; infection; therapy; ICAM-1; ss.
                                                                                                                                                                The invention relates to ebl-1 polypeptides that are encoded by the DBL (Duffy-binding like) gene family. The ebl-1 proteins are substantially identical to the Duffy-binding like) gene family. The ebl-1 proteins are substantially binding Protein (BABP) and Sialic Acid Binding Protein (BABP), which are soluble proteins that appear in the culture supernatant after erythrocytes infected with malaria release merozoites. Immunochemical studies indicate that DABP and SABP are the respective ligands for Plasmodium viox and Plasmodium falciparum Duffy and sialic acid receptors on erythrocytes. The ebl-1 polypeptides may be used to vaccinate against malaria, especially caused by P. falciparum. Immunization with the polypeptide provides effective protection against malaria. Sequences AAZ982397-304 represent primers used for isolating sequences encoding the conserved motifs of the DBL family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents an antisense inhibitor of ICAM-1. The invention relates to a pharmaceutical composition comprises an oligonucleotide (ON)
                                                    Isolated protein binding domains from Plasmodium vivax and Plasmodium falciparum erythrocyte binding proteins useful for vaccinating against malaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 55.6%; Pred. No. 9.3e+02; Matches 10; Conservative 7; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tsai YJ,
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 4 C; 5 G; 2 T; 0 U; 7 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ecker DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ICAM-1 antisense inhibitor, ISIS-1939.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 47; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1630 CCCAGCAGCAGCGCTG 1647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ||::|::|:||
||ccsmgsmgscagcagyTs 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cook PD,
                                                                                                                                  Example; Fig 3; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ48638 standard; DNA; 20 BP
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                  WPI; 2000-194198/17.
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admixed with a topical delivery agent. The compositions can be used for the delivery of a ribozyme, an external guide sequence, an antisense ON an antisense peptide nucleic acid, an aptamer or a molecular decoy. The ONS can be used to modulate expression of a cellular adhesion protein or modulate a rate of cellular proliferation. The compositions can also be used to treat psoriasis. They can also be used to treat e.g. lichen planus, toxic epidermal necrolysis, erythema multiforme, basal cell carcinoma, squamous cell carcinoma, malignant melanoma, Paget's disease, Kaposi's sarcoma, pulmonary fibrosis, Lyme disease and viral, fungal and bacterial infections of the skin. They can be used to treat humans and primates, avians including chickens and turkeys, domestic household, sport of farm annials including rate, mice, rabbits and quinea pigs, fish, reptiles and zoo animals. The compositions and methods may also be used to examine the function of various proteins and genes in vitro in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mutation in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a vector (I) comprising an isolated DNA sequence of a multiple tumour suppressor (MTS) gene having a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, multiple tumour suppressor; MTS; somatic mutation, cancer; diagnosis; germ line mutation; gene therapy; cytostatic; melanoma; leukaemia; astrocytoma; glioblastoma; lymphoma; glioma; Hodgkin's lymphoma; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New vector useful for gene therapy of cancer associated with mutumor suppressor gene, comprises DNA sequence of multiple tumor
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0
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Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cultured or preserved dermal tissues and in animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse P16 PCR primer SEQ ID NO:29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 12; Col 51; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAGAGGGAAGTGGTGGGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA39378 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94US-00215086.
94US-00215087.
94US-00227369.
94US-00251938.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 80.0
Matches 16; Conservative
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Example 3; Page 49; 116pp; English.

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polynucleotide sequence of the human MTSIE1-beta. (I) is useful for introducing wild-type MTS function to a cancerous or pre-cancerous cell which carries diminished or mutant MTS alleles for suppressing neoplastic growth of the recipient cells. (I) is also useful for increasing the recipient cells. (I) is also useful for increasing the level of expression of MTS gene even in tumour cells in which the mutant gene is expressed at a normal level but the gene product is not fully functional. A host cell transformed with (I) is useful as a model system to study cancer remission and drug treatment which promotes such cemission. The present invention relates to somatic mutations and germ line mutations in the MTS gene and their use in the diagnosis and prognosis of human cancer e.g. melanoma, leukaemia, astrocycoma, glioplastoma, lymphoma, glioma, Hodgkin's lymphoma, and cancers of the panceant invention ceptesents a PCR primer used in an example from the present invention
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Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;

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Gaps
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0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
rative 0; Mismatches 4; Indels
           Local Similarity 80.0 tes 16; Conservative
 Query Match
                         Matches
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AAC61834 standard; DNA; 20 BP. 06-MAR-2001 (first entry) RESULT 1268

Human, Fas; Apo-1; antisense compound; Fas ligand; Fap-1; hepatitis; Fas associated protein 1; protein tyrosine phosphatase; cancer; autoimmune disease; inflammatory disease; lymphoma; phosphorothioate; ss. Antisense oligonucleotide directed against human Fas ligand gene.

Homo sapiens. Synthetic

/note= "contains phosphorothioate linkages" ...5 *tag= a note= "2'-methoxyethoxy residues" /note= "2'-methoxyethoxy residues" Location/Qualifiers 16. .20 *tag= /*tag= modified base modified base misc_feature

WO200061150-A1

19-OCT-2000

10-APR-2000; 2000WO-US009540.

12-APR-1999; 99US-00290640. (ISIS-) ISIS PHARM INC.

Marcusson EG; Dean NM,

WPI; 2000-628395/60.

Antisense oligonucleotides for treating hepatitis and colon, liver or lung cancer are inhibitors of Fas, Fas ligand or Fas associated protein 1 (Fap-1) expression.

AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7740 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the numan genome, and in complex association studies and haphotyphing studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the agents and side effects from

Novel biallelic markers used to construct a high density disequilibrium

Claim 9; Page 2707; 2745pp; English.

map of the human genome.

Chumakov 1;

Cohen D, Blumenfeld M,

(GEST) GENSET

21-AFR-1998; 23-NOV-1998; 21-APR-1999;

WPI; 2000-013267/01.

99WO-IB000822. 98US-0082614P. 98US-0109732P.

Homo sapiens.

W09954500-A2

28-OCT-1999.

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                                                          AAC61821-39 represent antisense oligonucleotides which are directed against mucleid acids encoding human Fas ilgand. The specification describes antisense compounds which are targeted to the 5'-uutranslated region, translational start site, translational termination region or 3'-uutranslated region of nucleic acid molecules encoding Fas, Fas, Fagr-1 (Fas associated protein 1, protein tyrosine phosphatase). The antisense compounds are used to inhibit the expression of Fas, Fas, or Fapr-1 in cells or tissues. They are used to trast autoimmune or inflammatory diseases such as hepatitis. They can also be used to treat cancer, especially colon, liver or lung cancer or lymphoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human biallelic marker downstream amplification primer SEQ ID NO:11617.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            y Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 9.3e+02; hes 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 10 C; 3 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 cccrcrrcacareccacec 20
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ID AAZ77261 standard; DNA; 20
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8 \frac{1}{2} 
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Human, SHP-1; Src homology region 2-domain phosphatase; phosphorothioate, cytosolic tyrosine phosphatase; antisense oligonucleotide; cancer; leukaemia; inflammation; infection; ss.
                                                                                                                                                                                                                                                                                                                 Novel antisense oligonucleotides for modulating the expression of human SHP-1, especially for treating a disease or condition associated with SHP-1 expression, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to antisense oligonucleotides which modulate the expression of human SHP-1 (Src homology region 2-domain phosphatase) a cytosolic tyrosine phosphatase. The invention includes antisense molecules AAA09644-A09683 which have modified phosphorothioate internucleoside linkages which target various regions of the SHP-1 gene. The oligonucleotides inhibit the expression of human SHP-1 in cells or tissues, and may be used to treat diseases or conditions associated with SHP-1 expression e.g. cancers, specifically leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGP-3; tumour necrosis factor ligand; TNF ligand; Crohn's disease; type II transmembrane protein; B cell stimulatory factor; inflammatory disorder; immune disorder; rheumatoid arthritis; lupus and graft versus host disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer for murine cDNA encoding an AGP-3 polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          210 GCAGATAGGCCTGGATGAGA 229
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                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Col 41; 33pp; English.
                                                                                                                                                               99US-00358685.
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99US-0166271P.
                                                                                                                                                                                           9905-00358685.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                     Bennett CF, Cowsert LM;
                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                   WPI; 2000-593714/56.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200047740-A2.
                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-DEC-2000
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                                                                                                                                                               21-JUL-1999;
                                                                                                                                                                                           21-JUL-1999;
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                                                                                                    US6121047-A.
                                                                                                                                19-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA63936;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus sp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to von Willebrand factor. The present sequence represents a primer used in the methods of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Method for purifying factor VIII using chimera antibody to von Willebrand
pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the present invention
                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kim BJ;
                                                                                                              Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        / Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 65.0%; Pred. No. 9.3e+02; Nes 13; Conservative 4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                              Primer #13 in invention relating to von Willebrand factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lee JM,
                                                                                Sequence 20 BP; 9 A; 0 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 3 C; 5 G; 3 T; 0 U; 4 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human SHP-1 antisense oligonucleotide SEQ ID 31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oh HG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lee JS,
                                                                                                                                                                         1237 CACTICATCITCCGTAICTI 1256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      140 AGATCAAACGGCAGCTGTCA 159
                                                                                                                                                                                                                                                                                                                                                                                                        Von Willebrand factor; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 4; 12pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | | |::|:| ||||| AGGTSMARCTGCAGSAGTCA 20
                                                                                                                                                                                             20 CTCTCCCTCTTCCATATCTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GREC ) KOREA GREEN CROSS CORP.
                                                                                                                                                                                                                                                                                   AAS14488 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98KR-00002265.
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                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                  AAS14488;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     factor.
                                                                                                                                                                                                                                                       1270
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Page 604

schultz621-3.rng

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PCR primers AAA61936-37 were used to amplify cDNA encoding a murine AGP-3 polypeptide. AGP-3 is a tumour necrosis factor (TNF) ligand family member. AGP-3 is a type II transmembrane protein, and is a potent B cell stimulatory factor. Expression of AGP-3 correlates to increases in the number of B cells and immunoglobulins produced. AGP-3 proteins, antibodies, and nucleic acids may be used to treat inflammatory and immuno disorders, e.g. rheumatorid arthritis, Crohn's disease, lupus and graft versus host disease. The nucleic acids may be used to regulate the expression of an AGP-3 related protein. The AGP-3 proteins, antibodies and nucleic ands are also useful for the detection of AGP-3 agonists, antagonists and characterizing interactions with AGP-3 related proteins
                                                                            Novel polypeptides comprising tumor necrosis factor ligand family protein; useful for treating inflammatory and immune disorders, e.g. rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cook PD, Tillman L, Hardee GE, Ecker DJ, Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ICAM-1; cellular adhesion; expression; modulation; antisense; non-parenteral; delivery; uptake; administration; emulaton; ulcerative colitis; Crohu's disease; inflammatory bowel disease; cellular proliferation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ICAM-1 targetted phosphorothioate oligonucleotide ISIS 1939.
                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                              Seguence 20 BP; 6 A; 7 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                             Disclosure; Page 36; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            916 CTGTTCCTGTTCCAGCTGCT 935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
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                                                WPI, 2000-558217/51.
                Boyle WJ, Hsu H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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New oligonucleotide compositions used for the non-parenteral delivery of

WPI; 2000-072428/06.

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This invention describes a novel method for determining the identity of a specific nucleotide at one or more defined sites in a target nucleic acid polymer involves formation of a detectable primer extension product if the specific nucleotide is present at the defined site in the target nucleic acid. The method is specifically used to detect point mutations which are associated with altered pathogenicity or resistance to therapy
                                                                                                Sequences AAZ49336-Z49343 and AAZ49390 represent antisense oligonucleotides designed to modulate cellular adhesion. The invention relates to new compositions for the non-parenteral delivery of oligonucleotides comprising at least one oligonucleotide in an emulsion. Oligonucleotides delivered via the compositions of the invention can be used to modulate expression of a cellular adhesion protein, modulate a rate of cellular proliferation, or have biological activity against evikaryotic pathogens or retrovituses. They can be used for treating conditions including e.g., ulcerative colitis, Crohn's disease, inflammatory bowel disease or undue cellular proliferation. The compositions can enhance the local and systemic uptake and delivery of nucleic acids via non-parenteral routes of administration (e.g., via the alimentary canal, skin, eyes, pulmonary tract, urethra or vagina)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying the nucleotide at specific position in a target sequence for detecting disease-related point mutations involves extending a primer that binds adjacent to the specific site to incorporate a labeled deoxynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection, primer extension, point mutation, pathogenicity, therapy; cancer, genetic disease, K-ras, human, PCR primer, mutation, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gape
e.g. antisense oligos, ribozymes, peptide nucleic acids, molecular decoys, external guide sequences or aptamers.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 7; Col 17-18; 14pp; English.
                                                                 Claim 80; Page 37; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 chchecechaerecreece 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ44889 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human K-ras PCR primer R6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-146544/13.
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in a microorganism, particularly human immune deficiency virus or with cancer or a genetic disease (or susceptibility to it) in humans, but more generally can be used to detect mutations in RNA or DNA from animals, plants or microorganisms. By selecting a primer that binds adjacent to the specific site, variations at this site can be detected following incorporation of only a single dNTP. The method requires only a few, simple manipulations, making is suitable for routine use, and allows quantification of the proportion of mutated calls in a mixed population, down to 0.5% of this population. The method is easily automated. This sequence represents a PCR primer used to detect a mutation in the human K -ras gene

Sequence 20 BP; 3 A; 3 C; 7 G; 7 T; 0 U; 0 Other;

ö 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels 1311 GACATACAACTACCCCAAGT 1330 Matches 16; Conservative Similarity Query Match Local

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Gaps

20 GAGCTCCAACTACCACAAGT 1 셤 Š

AAZ89211 standard; DNA; 20 BP. AAZ89211; RESULT 1275

(first entry) 09-JUN-2000 Human glyceraldehyde-3-phosphate dehydrogenase forward PCR primer.

Human; expression profile, Three Prime End Amplification; TPEA; glyceraldehyde-3-phosphate dehydrogenase; PCR primer; ss.

Homo sapiens.

WO200008208-A2.

17-FEB-2000

99WO-GB002579. 05-AUG-1999;

98GB-00017055. 05-AUG-1998;

(MEDI-) MEDICAL RES COUNCIL.

Freeman TC, Richardson PJ, Dixon AK;

WPI; 2000-224033/19.

Reverse transcription of mRNA species used for expression profiling of single cells by employing a first heeled primer to provide first strand cDNA species and then a second heeled primer population to generate second strand cDNAs.

Example 1; Page 29; SOpp; English.

This invention describes a novel process (M1) of reverse transcribing mRNA species present in a sample from an organism by: (a) reverse transcribing the mRNA species using a first heeled primer, to provide a first strand CDNA species using a second heeled primers being second CDNA species using a second heeled primers being such that the reverse heel portions of the second heeled primers being such that the reverse transcribed first strand CDNA species are capable of hybridizing to at least one second primer. The processes can be used for expression profiling of single cells. The polynucleotide comprising an oligo d(T) sequence and a heel sequence 5, can be used for the reverse transcription of mRNA species in a sample. The polynucleotide primer population of clam (4) can be used for the synthesis of second strand CDNA from a population of first strand CDNA species. Single cell CDNA libraries can be made for subsequent detailed analysis of gene expression and the AAZB92111/ AAZB92111/ AAZBAAC AAZB XXX AAC AAZB XXX Huma XXX Huma XXX HODE XXX HODE XXX WO20 XXX

discovery of novel genes. Small samples can be used and allow the utilization of the large amount of sequence data available for further understanding of disease processes and repertance of complex issues. The invention provides a rapid, robust and reproducible procedure (aslee Three Prime End Amplification (TPEA). Optionally with PCR (TPEA-CR). Prior art methods for the analysis of gene expression within single cells or small tissue samples are limiting. Whilst in situ hybridization techniques provide detailed information about the cellular expression pattern of a gene in intact tissue the technique is laborious and unable to analyze multiple transcripts in a single preparation. The methods presented in the disclosure provide a more straightforward, reproducible and reliable cDNA amplification procedure for small mRNA samples where expression profiling can be conducted. The amplification technique can be expression profiling can be conducted. The amplification technique can be carried out in a single tube with a need for only limited manual arge numbers of samples can be analyzed. There is a intervention and large numbers of samples can be analyzed. There we bias towards more uniform length cDNA molecules ensuring that even the calatively low abundance mRNA species are transcribed and optionally amplified at the same level of efficiency as more abundant mRNA species. Alzapalol 289253 represent the primers described in the method of the invention

Sequence 20 BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;

Gaps ö Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02; 0; Mismatches 4; Indels 0.8%; 16; Conservative Local Similarity Query Match Matches

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RESULT 1276 AAA11188

AAA11188 standard; DNA; 20 BP. AAA11188;

11-OCT-2000 (first entry)

Mouse multiple tumour suppressor 1 Elbeta p16-specific reverse primer.

Variant; human; multiple tumour suppressor; MTS; mutation; melanoma; cancer; diagnosis; PCR primer; ss.

Mus sp.

US6037462-A.

98US-00120130 22-JUL-1998; 14-MAR-2000

94US-00215086. 94US-00215087. 94US-00227369. 94US-00251938. 94US-00214582, 18-MAR-1994; 18-MAR-1994; 18-MAR-1994

95US-00480810 95WO-US003316 14-APR-1994; 01-JUN-1994; 17-MAR-1995; 07-JUN-1995;

(MYRI-) MYRIAD GENETICS INC.

Kamb A;

WPI; 2000-269915/23.

New mutants of the human multiple tumor suppressor gene, useful as diagnostic markers of cancer, contain specific base alterations or deletions.

Example 12; Col 50; 72pp; English.

The invention relates to variants (AAA11196-A11206) of the human multiple potential anticancer agents. This sequence screen for the mouse MTS1 Elbeta sequence 3ennett CF, Mirabelli CK, Baker BF; Example 10; Page 176; 199pp; English. 505 GAGGGCTACCTGGAGAAGCT 524 20 GAAGGCTTCCTGGACACGCT 1 AAZ48909 standard; DNA; 20 BP 98US-00085759 99WO-US011548 29-MAR-2000 (first entry) Local Similarity 80.0 ies 16; Conservative (ISIS-) ISIS PHARM INC. WPI; 2000-072600/06. Homo sapiens. WO9961462-A1. 26-MAY-1999; 27-MAY-1998; 02-DEC-1999 AAZ48909; Query Match Best Loca Matches AAZ48909, d 8ò

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tumour suppressor 1 (MTS1) gene (AAA11165). The variants have the following changes relative to this sequence: A at any of positions 265, 442, 330 and 329; T at any of positions 172, 238, 341 and 148 and deletions of mucleoxides 290-294, 172-179 or 128-129. The variants are somatic mutations of MST1, indicative of predisposition to melanowa and many other cancers, so detecting them is useful for diagnosis, prognosis and monitoring of cancer (including prenatal analysis). Cells and animals that express the variants are useful as model systems for identifying potential anticancer agents. This sequence represents a primer used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
rative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
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Human ICAM-1 antisense inhibitor, ISIS #1939.
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Antisense inhibitor; human; ICAM-1; intercellular adhesion molecule-1; vascular cell adhesion molecule-1; hyperproliferative disorder; VCAM-1; emodothelial leukocyte adhesion molecule-1; ELM-1; skin condition; cancer; viral infection; tumour; diappedesis; graft versus host disease; arthritis; infection; autoimmune disorder; multiple sclerosis; stroke; purchile diabetees mellitus; arthritis; myasthenia gravis; therapy; pemphigus vulgaris; systemic lupus erythematosus; autoe myocarditis; cardiovascular disorder; dilated cardiomyopathy; ischaemic heart disease;

New antisense oligonucleotides, used for treating e.g. inflammatory conditions, psoriasis, graft rejection, cancers, infections, cardiovascular disorders or autoimmune disorders.

This sequence is an antisense oligonuclectide of the invention. The antisense oligonuclectides are targeted to a nucleic acid encoding a cellular adhesion molecule (CAM) and is capable of modulating the expression of the CAM. They particularly inhibit intercellular adhesion molecule-1 (ICAM-1), vascular cell adhesion molecule-1 (UCAM-1), or endotherial leukocyte adhesion molecule (ILAM-1). The antisense oligonuclectides can be used to modulate CAM activity in mediating cell:cell interactions and subsequent cellular and biological responses,

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antisense sequences can be used for modulating the synthesis of a CAM.

They can be used for modulating the synthesis of a CAM.

They can be used for treating an animal suspected of having or being proceed to a disease or condition associated with a CAM. Oligonucleotides targeted to ICAM-1 can be used for treating an inflammatory disease or condition e.g. inflammatory bowel disease such as Crohr's disease or condition e.g. inflammatory bowel disease such as Crohr's disease or condition e.g. inflammatory bowel disease such as Crohr's disease or condition e.g. inflammatory bowel disease such as CAM-1 sequences or intermatitis, rhemmatoid arthritis, allograft rejection, cancer, or conting cor reducing corticosteroid use in a patient or for reducing cyclosporine use in a patient. The oligonucleotides can also be used for reducing corticosteroid use in a patient or for reducing cyclosporine use in a patient. The oligonucleotides can also be used for detection and diagnosis. They can also be used for treating e.g. hyperpolifestative disorders, tumours, diapedesis, graft versus host disease, arthritis, infections, autoimmune disorders, e.g. autoimmune to some of juvenile disorders mellitus, mysathenia gravis, pemphigus culgaris, systemic lupus erythematosus, cardiovascular disorders, myocardial ischaemia/reperfusion injury, dilated cardiomyopathy, acute myocarditis, ischaemic heart disease or stroke
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Typing polymorphic genes, useful to assess the association of alleles with diseases and in disease diagnosis, uses a taxonomy based sequence analysis in which a typing tree based on distinguishing sequences is constructed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides a novel method for typing genes, particularly human leukocyte antigen (HLA) coding sequences. The methouses DNA sequences and a taxonomy-based sequence analysis method to assign alleles for HLA-DQA1, HLA-DQB1 and HLA-DRB. These alleles have been linked to diseases such as diabetes, IgA deficiency, multiple sclerosis, cancer, clinical and immunological manifestations of HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
Live 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 64; 125pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      226 GAGAGTGGTGGTGGTGG 245
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Best Local Similarity 80.0
Matches 16; Conservative
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infection, coeliac disease, idiopathic nephrotic syndrome, immune

BP.

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AAA94747 standard; DNA; 20
                                   AAA94747;
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  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides a novel method for typing genes, particularly human leukocyte antigen (HLA) coding sequences. The method uses DNA sequences and a taxonomy-based sequence analysis method to assign alleles for HLA-DQA! HLA-DQBI and HLA-DRB. These alleles have been linked to diseases such as diabetes, IgA deficiency, multiple sclerosis, cancer, clinical and immunological manifestations of HIV infection, coeliac disease, idiopathic nephrotic syndrome, immune responses to parasite antigens, pemphigus vulgaris, inflammatory bowel disease, rheumatoid arthritis, allergy and other inflammatory diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Typing polymorphic genes, useful to assess the association of alleles with diseases and in disease diagnosis, uses a taxonomy based sequence analysis in which a typing tree based on distinguishing sequences is
           responses to parasite antigens, pemphigus vulgaris, inflammatory bowel disease, rheumatoid arthritis, allergy and other inflammatory diseases
                                                                                         Gaps
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                                                                 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ve 0; Mismatches 4; Indels
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                                             Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Brunham K;
                                                                                                              1427 TCTCCGCAGAGGATGCCATG 1446
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                                                                                                                                                                                                                                                            Gene typing PCR primer SEQ ID NO:
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                                                                                                                                   1 rcccccccacacarrccrc 20
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                                                                                        Conservative
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                                                                          Local Similarity
nes 16; Conserv
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Local Sim
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                                                                                                                                                                     RESULT 1279
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The present invention relates to an aggregated composition comprising a polypeptide having the transport function of herpesviral transport protein VP22. The aggregates can be useful for delivery of oligonucleotides and proteins into cells. The present sequence is one such oligonucleotide which may be delivered into cells using the method of the present invention. The aggregated composition is useful for preparing medicament for therapy or prophylaxis of a disease and for delivering molecules to cells in vitro. The aggregates are delivered to target cells such as tumour cells in vitvo and are useful for treating pooriasis, eczema or skin cancer
                                                                                                                                                                                                                                                                                            /mod_base= OTHER /mod_base are phosphorothioate deoxymucleotides /note= "All bases are phosphorothioate deoxymucleotides Optionally labelled at 3' end with fluorescein or at 5' end with blottin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Aggregated composition suitable for phototherapy or prophylaxis of psoriasis, eczema or skin cancer and for delivering nucleic acids and proteins into cells, comprises transport protein VP22 and an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                       VP22; gene therapy; tumour; psoriasis; eczema; skin cancer;
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                                                                                                                                                                                                               Location/Qualifiers
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nes 16; Conservative
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                                                   Oligonucleotide #1.
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modified_base
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19-JAN-2001
                                                                                                                                                              Unidentified
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RESULT 1280

Human c-raf kinase antisense oligonucleotide #11 (Isis #5149).

94US-00250856. 95WO-US007111. 96US-00756806.

98US-00143214.

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c-raf is a serine-threonine-specific protein kinase and is thought to play a fundamental role in signal transduction, and cell proliferation control. The present sequence is an antisense oligomuclectide. This sequence is targeted to human c-raf gene, resulting in c-raf expression inhibition. The present sequence may be useful for treating and rafsociated cell hyperproliferation conditions such as cancer, hyperplasias, pulmonary fibrosis, angiogenesis, psoriasis, and smooth muscle cell proliferation in blood vessels etheroselerosis and smooth muscle cell proliferation in blood vessels e.g. stenosis or restenosis following angioplasty. Also, the present sequence may be useful for treating inflammatory disorders such as tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Interleukin 1 receptor antagonist short tandem repeat primer SEQ ID NO:7.
                          Human, c-raf; protein kinase; antisense oligonucleotide; cancer; signal transduction; hyperplasia; pulmonary fibrosis; angiogenesis; psoriasis; atherosclerosis; mosch muscle cell proliferation; stenosis; restenosis; inflammatory disorder; tissue graft rejection; endotoxin shock; glomerular nephritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Short tandem repeat; primer; STR; susceptibility; HIV; infection; AIDS; detection; polymorphism; interleukin 10 promoter; IL-10; chromosome position 2q12; interleukin 1 receptor antagonist; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense oligonuclectides targeted to nucleic acid molecule encoding human raf useful for diagnosis, treatment of raf-associated cell proliferative conditions such as cancer, psoriasis or blood vessel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence may be useful for treating inflammatory disorders
graft rejection, endotoxin shock and glomerular nephritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                              Boggs RT, Monia BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-531424/48.
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31-MAY-1995;
26-NOV-1996;
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                                                                                                                                                  Homo sapiens
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The present invention describes a method for predicting susceptibility to HIV infection or HIV progression in a subject. The method involves detecting a polymorphism in a human interleukin-10 (Li-10) promorer, where the presence of the polymorphism interleukin-10 (Li-10) promorer, infection or HIV progression. The method provides prognostic information to presence infected with HIV virus and is useful to help select treatments (such as administration of II-10 or gene therapy with II-10). The presence of polymorphism is useful as predictor that very aggressive treatment could substantially eradicate the virus from the infected person. The method is useful for the generation of normograms or other predictive algorithms that can be used, in association with allele status, to prognose probable survival or years to development of AIDS following HIV seroconversion. It indicates that increased expression of the II-10 gene helps to reduce HIV-1 infection and pathogenic progression and bushoes a variety of new therapeutic interventions in the treatment of HIV and all and and and pathogenic progression and bushoes a variety of new therapeutic interventions in the treatment of HIV and and and pathogenic progression when the predictive and
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                                                                                                                                                                                                                                                                                                                                Predicting susceptibility to HIV infection or progression useful for selection of therapeutic treatment for persons infected with HIV virus, comprises detecting polymorphism in human interleukin-10 promoter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of HIV disease. The present sequence represents a short tandem repeat (STR) primer which is used in an example from the present invention
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80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
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                                                                                                                                                                              (USSH ) US DEPT HEALTH & HUMAN SERVICES.
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                                                                                                                                                                                                                                    Smith MW, Shin HD, O'brien SJ;
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                                                                       06-APR-2000; 2000WO-US009355
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Best Local Similarity 80.0
Matches 16; Conservative
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                                                                                                                             09-APR-1999;
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Gaps ô

(first entry)

WO200061811-A2

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Sun Y;

Habben JE,

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Polymucleotide sequences AAC83101 - AAC83113 encode proteins AAB35794 -
AAB35806 which are involved in regulating the cell cycle. The protein and
DNA sequences have been isolated from Zea mays (corn), and the invention
also includes oligomucleotides. The cell cycle polymucleotide sequences
are useful for producing transgenic plants such as maize, soybean,
cunflower, sorghum, canola, wheat, alfalfa, cotton, rice, barley and
milet with increased levels of cell cycle gene activity, such as
activity of cyclin and cyclin-dependent kinases. The DNA sequences are
also useful as probes for detecting deficiencies in the level of mRNA in
screening for desired transgenic plants, for detecting mutations in the
gene, for monitoring upregallation of expression or changes in enzyme
activity in screening assays of compounds, for detecting any number of
allelic variants orthologs or paralogs of the gene, and site-directed
mutagenesis in eukaryotic cells. The DNA sequences are also useful for
recombinant expression of the encoded polypeptides and as immunogens for
preparing and screening antibodies. A transgenic plant comprising an
copiession cassette including a cell cycle regulatory gene is useful for
assaying enzyme agonists and antagonists, and as immunogens or antigens
to obtain antibodies. The antibodies are useful in assaying expression
clevels of cell cycle regulatory proteins for identifying and isolating
mucleic acids from other species, and for purification of the proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide; p38 mitogen activated protein kinase; MAPK; antirheumatic; antiarthritic; immunosuppressive; cardiant; heart disease; antiinflammatory; autoimmune disease; rheumatoid arthritis; apoptosis; phosphorothioate; ss.
                                                                                   Nucleic acids useful for producing transgenic plants, preferably maize, with increased cell cycle gene activity, preferably activity of cyclin and/or cyclin-dependent kinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 13.6; DB 1; Length 20; ilarity 80.0%; Pred. No. 9.3e+02; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine p38beta antisense oligonucleotide SEQ ID 75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 2 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                           Disclosure; Page 118; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           279 ICCIGGGGAACTICGIICIG 298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 TCAAGGGAAATTGGTTCTG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC79550 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-APR-2000; 2000WO-US008794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                         WPI; 2000-687333/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 16; Conserv
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                   Helentjaris TG,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC79550;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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99US-00286904

(first entry)

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This invention relates to antisense compounds 8-30 nucleobases in length targeted to the 5'-untranslated region, translational start alte, translational start alte, translational termination region or 3'-untranslated region of a nucleic acid encoding a p18 mitogen activated protein kinase (MAPK), where the antisense oligonuclectides inhibit the expression of MAPK. Sequences AAC79501 represent human p18alpha MAPK and p18beta MAPK cDNA sequences. AAC79510 represent human p18alpha antisense oligonuclectides, while AAC79510 represent human p18alpha antisense oligonuclectides, while AAC79521 and AAC79521 and AAC79510 represent human p18alpha cDNA sequence AAC79523 and antisense oligonuclectides AAC79523 - AAC79531 and antisense oligonuclectides AAC79531 - AAC79537 and antisense oligonuclectides have antishense oligonuclectides are useful for inhibiting the expression of p18 MAPK in cells or tissues. The oligonuclectides are useful for inhibiting the expression of p18 mapk in a think hidseases such as inflammatory or autoimmune diseases e.g. rheumatoid arthitis, or heart diseases. The oligonuclectides are also useful for arthitis are also useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NPRS, non-ribosomal peptide synthetase; adenylation domain; A domain; PCR primer; antibiotic; immunosuppressant; cytostatic; antiviral; antihelminthic; fungicidal; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                             protein kinase
and treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.6; DB 1; Length 20; 30.0%; Pred. No. 9.3e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  B. brevis NRPS gene A domain PCR primer SEQ ID No: 10.
                                                                           Antisense compound targeted to p38 mitogen activated inhibits protein kinase and is useful for diagnosing inflammatory, autoimmune and heart disease.
Popoff I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 6 A; 7 C; 5 G; 2 T; 0 U; 0 Other;
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  Mckay R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Marahiel MA, Stachelhaus T, Mootz H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inhibiting inflammation or apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1153 GACATGTGGGGTGTGGCCTG 1172
                                                                                                                                                                    Example 5; Page 54; 90pp; English.
  Gaarde WA, Nero PS,
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match
Best Local Similarity 80.0°
Matches 16, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MARA/) MARAHIEL M A
(STAC/) STACHELHAUS
(MOOT/) MOOTZ H.
                                          WPI; 2000-664982/64.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-SEP-2003
26-JAN-2001
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  Monia BP,
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(MOOT/)
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The present invention describes a method of modulating bone resorption comprising administering leptin or a derivative under conditions suitable for the modulation of osteoclastogenesis. This is useful in the treatment of osteoporosis and paget's disease. No further information about this
                                                                                                                                                                                                                                                                                                             This invention describes a novel method for the targeted non-ribosomal synthesis of peptides (I) of required structure, comprising altering one or more A (adenylation) domain-encoding DNA segments (II) that encodes a non-ribosomal peptide synthetase so that the expression product of the altered (II) can produce (I), is new. Alterations in the A-domains are made according to a non-ribosomal code reproduced in the Specification. The method is used to synthesize (I) with antibiotic, immunosuppressant, cytostatic, antiviral, antihelminthic, fungicidal or surface-active properties, and to alter specificity and/or activity of known piologically active compounds, e.g. to improve thair solubility by replacing hydrophobic amino acids with hydrophilic ones, or vice verse. AAA97960-A97995 represent PCR primers used to illustrate the method of the invention. (Updated on 15-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modulating bone resorption in human or animal for treating osteoporosis or Paget's disease, comprises administering leptin, its derivative, homologue, analog, chemical equivalent, antagonist or agonist.
                                                                                                Non-ribosomal synthesis of peptides, e.g. antibiotics or immunosuppressants, using non-ribosomal peptide synthase with targeted modifications in adenylation domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bone resorption modulation; leptin; osteoporosis; Paget's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bone resorption modulation method related sequence SEQ ID NO: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1515 ACTAAAGGAGATTCAGCTAC 1534
                                                                                                                                                                                                                                                             Example 1; Page 39; 52pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 ACTACAGCAGGCTCAGCTAC 20
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                                    WPI; 2000-572182/53.
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                                                                                                                                                                                                                 Human, glycogen synthase kinase 3 alpha, antidiabetic, cytostatic, antisense therapy, diabetes; hyperproliferative disorder, inflammation, neurological disorder, tumour, haematopoietic disorder, infection; hyperproliferative disorder; developmental disorder; antisense;
                                                     Gaps
                                                                                                                                                                                               Human glycogen synthase kinase 3 alpha antisense oligo ISIS #116602.
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                                 Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                           note = "Phosphorothioate backbone"
                 Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
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/note= "Methoxyethyl residues"
16
                                                                                                                                                                                                                                                                                                                                                                                note = "Methoxyethyl residues"
sequence is given in the specification
                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                         122 CCATGGATCGGATGAAGAAG 141
                                                                                                                                                                                                                                                                                                                 *tag= a
mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                        'mod_base= OTHER
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16..20
/*tag= c
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/mod_base= m5c
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mod_base= m5c
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                                   Query Match
Best Local Similarity 80.0%;
Matches 16; Conservative
                                                                                                                                         AAD14761 standard; DNA; 20
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                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                             AAD14761;
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                                                                                                                         RESULT 1287
AAD14761
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RESULT 12
AAC81175/
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/mod_base= OTHER
/note= "OTHER = F1 (CH2)6-PO-thymine, where F1 is flavine
                                                                                                                                                                                                                                                                                                 The invention relates to an antisense compound 8 to 30 nucleobases in length targetted to a nucleic acid encoding glycogen synthase kinase 3 alpha. The antisense compound specifically hybridises with and inhibits the expression of glycogen synthase kinase 3 alpha. The antisense compound is useful for the treatment of a diseases associated with glycogen synthase kinase 3 alpha such as diabetes, a neurological disorder, a haematopoietic disorder, a hyperproliferative disorder or a developmental disorder. The antisense compounds may also be used prophylactically to prevent or delay infection, inflammation or tumour formation. The present sequence is a phosphorothioate antisense oligonucleotide targetted to human glycogen synthase kinase 3 alpha DNA
                                                                                                                                                                                                                          Antisense compound 8 to 30 nucleobases in length comprising a compound that is targeted to a nucleic acid molecule encoding glycogen synthase kinase 3 alpha, useful for the treatment of e.g. diabetes and hyperproliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analytical support; genomic sequencing; mutation detection;
pharmaceutical development; 8s.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 12 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                           Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                  Example 15; Page 82; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   556 CTCAGCCGCCGCCTCGTCG 575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 crcccrccrccrccrccccc 20
                                                                                                                                                                                           Butler MM,
 /*tag= m
/mod_base= m5c
                                          mSc
                                                           /*tag= o
/mod_base= m5c
                                                                                                                                16-JAN-2001; 2001WO-US001411.
                                                                                                                                                  21-JAN-2000; 2000US-00488856
                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%;
Similarity 80.0%;
                               /*tag= n
/mod_base= r
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                      (ISIS-) ISIS PHARM INC
                                                                                                                                                                                           Mckay R,
                                                                                                                                                                                                              WPI; 2001-442247/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide #7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
modified_base
                                                                                          WO200152865-A1
                      modified base
                                                   modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                               Ma.
Local Si...
16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-FEB-2002
                                                                                                             26-JUL-2001
                                                                                                                                                                                           Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABA04587;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1288
ABA04587/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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The present invention relates to an analytical support, to which a number of oligonucleotides are fixed. The oligonucleotides are labelled with a fluorescent compound, the fluorescence of which varies when the oligonucleotide hybridises to its complement. The analytical support is useful in hybridisation testing for identification of specific nucleic acids, such as genomic sequencing, detecting mutations or pharmaceutical development. The present oligonucleotide was used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Support for hybridization analysis of nucleic acids for sequencing techniques, comprises an array of oligonucleotides having a label where the fluorescence changes follow hybridization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human bcl-6 phosphorothioate antisense oligonucleotide, SEQ ID NO:41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, bcl-6; transcriptional repressor; germinal centre formation, Th-2 mediated antibody affinity maturation, apoptosis regulator; chromosome 3q27; lymphoma, acute lymphoblastic leukaemia; post-transplant lymphoproliferative disorder; expression inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                            Fontecave M, Decout JL,
and PO is a phosphate group"
                                                                                                                                                                                                                                                                                     (COMS ) COMMISSARIAT ENERGIE ATOMIQUE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 8; Page 18; 33pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 GAGAGGGAAGTGGTGGGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%;
                                                                                                                                                                    23-FEB-2000; 2000FR-00002236.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAC81175 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-048959/06
                                                                                                                                                                                                                                                                                                                                            Peltie P,
                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-628265/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-OCT-1999;
                                                         FR2805348-A1
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                                                                                                             24-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                            Cuzin M,
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sequences AAC81144-C81223 represent antisense oligonucleotides targetted to the human bcl-6 gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human bcl-6 mRNA, and were analysed for their effect on bcl-6 mRNA levels by quantitative real-time PCR. Bcl-6 (also known as B-cell CLL/ lymphoma 6, zinc finger protein 51 and iLAZ3) is a sequence-specific DNA-binding transcriptional represers. The bcl-6 gene is expressed in germinal centre B- and T- cells and is required for germinal centre formation and Th-2 mediated antibody affinity maturation. Bcl-6 may also play a role in the region which undergoes a high frequency of translocation events. Such chromosomal translocations can result in aberrant forms of Bcl-6, which are strongly implicated in the pathogenesis of several types of lymphoma, and have also been reported in acute lymphoblastic leukaemia and post-transplant lymphoproliferative disorders. The oligonucleotides of the cinvention are useful for diagnosis, prevention and treatment of conditions associated with aberrant forms of bcl-6, such as lymphomas, acute lymphoblastic leukaemia and post-transplant lymphoproliferative disorders.
           Antisense compounds which specifically hybridize with and inhibit human bcl-6 expression, useful for treating bcl-6 related disorders, and preventing or delaying inflammation or tumor formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel multiple tumor suppressor gene useful for diagnosing, prognosing and treating cancers, such as melanoma, leukemia, glioblastoma and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; multiple tumour suppressor; MTS; cancer; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.6; DB 1; Length 20; 30.0%; Pred. No. 9.3e+02; e. 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 20 BP; 4 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                    Claim 14; Col 41-42; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           494 TCCGGCTGCCTGAGGGCTAC 513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 rccedarccrereredacaac 1
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94US-00215087.
94US-00251938.
95WO-US003316.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF58196/c
ID AAF58196 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAR-1994;
18-MAR-1994;
01-JUN-1994;
17-MAR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Primer #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF58196;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1290
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Matches
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Human; cot oncogene; antisense therapy; inflammation; cancer; antisense; immune system disorder; prophylaxis; cytostatic; immunomodulator; Tpl-2; est; phosphorothioate backbone; ss.
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                                                                                                   The present invention relates to human multiple tumor suppressor-2 (Mgene. The invention is useful for diagnosing, prognosing and treating cancers. It is also useful for screening drugs for cancer therapy and
                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cot oncogene antisense oligonucleotide, ISIS 116381.
                                                                                                                                                                                                                                                                                                                                                        4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16. 720
/ *tag= c
//md base= OTHER
//note= "2'-methoxyethyl residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mod_base= OTHER
note= "2'-methoxyethyl residues"
                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                              0.8%; Score 13.6; DB 1;
80.0%; Pred. No. 9.3e+02;
vative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                       Example 12; Col 50; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                             505 GAGGCTACCTGGAGAGCT 524
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/mod_base=
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                                                                                                                                                                                                                                                                                        Ouery Match
Best Local Similarity 80.07
Marches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-463936/50.
Hodgkin's lymphoma.
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modified_base
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Synthetic.
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                                                                                                                                                                                                gene therapy
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(ISIS-) ISIS PHARM INC. Popoff I, Cowsert LM;

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The invention relates to antisense oligonucleotides, compositions and methods for modulating cot oncogene expression. The cot oncogene is also known as Tpl-2 and est. The compositions comprise antisense compounds, particularly antisense oligonucleotides, targetted to nucleic acids encoding cot oncogene. The antisense oligonucleotides are useful for associated with expression of cot oncogene and for treating diseases associated with expression of cot oncogene, e.g. inflammation, cancer of disorders of the immune system. The antisense oligonucleotides are also useful for diagnosis or prophylaxis or as research reagents and kits. The present sequence is human cot oncogene antisense oligonucleotide, ISIS 116381. This sequence was targetted towards the coding region of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, ss; PARP; Poly (ADP-ribose) polymerase; antisense oligonucleotide; cytostatic; nootropic; neuroprotective; antiinflammatory; antidiabetic; immunosuppressant; hyperproliferative disorder; cancer; cellular injury; oxidative stress; neurological disorder; parkinsonism; apoptoais; meningitis-associated intracranial complication; ischaemia; probe; inflammatory disorder; autoimmune disorder; arthritis; diabetes.
expression of cot oncogene, such as inflammation, cancer or immune system disorders.
                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mod_base= OTHER
note= "All cytidine residues are 5-methyl cytidine"
                                                                                                                                                                                                                                                                                                                                                                                         .
                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= c
/mod_base= OTHER
/mode= "2'-methoxyethyl nucleotides"
16.20
/*tag= d
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "2' methoxyethyl nucleotides"
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*tag= a "mod base= OTHER"
'note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 5 A; 2 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PARP-3 antisense inhibitor ISIS #126059.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                  Example 15; Col 41; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                          627 GGACAACTGGGCGAGGGTA 646
                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-MAR-2000; 2000US-00517467.
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*tag≈ b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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modified base
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                                                                                                                                                                                                                                                                                          cot oncogene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS45859;
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                                                                                                                                                                                        The invention relates to antisense oligonuclectides targeted to human PARP nucleic acid and inhibiting expression of human PARP. PARP (Poly M. (A.P.-tibose) polymerase plays an important role in chromatin decondensation, but a personation, but a discondensation, but a differentiation and apoptosis. The antisense oligonucleotide inhibitors are useful for inhibiting the expression of PARP in human cells or tissues. They are also useful for treating a human with a disease associated with PARP especially hyperproliferative disorders (e.g. cancer), cellular injury resulting from oxidative stress, neurological (e.g parkinsonism, meningitis-associated intracranial complications and ischaemia), inflammatory and autoimmune disorders (e.g architits) and diabetes. The present sequence is an antisense oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ss; PARP; Poly (ADP-ribose) polymerase; antisense oligonucleotide; cytostatic; nootropic; neuroprotective; antiinflammatory; antiidiabetic; immunosuppressant; hyperproliferative disorder; cancer; cellular injury; oxidative stress; neurological disorder; parkinsonism; apoptosis; meningitis-associated intracranial complication; ischaemia; probe; inflammatory disorder; autoimmune disorder; arthritis; diabetes.
                                                                                                            Antisense compound useful for treating hyperproliferative, neurological, inflammatory and autoimmune disorders and diabetes inhibits human PARP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mod_base= OTHER
/note= "All cytidine residues are 5-methyl cytidine"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
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/*tag= c
/mod_base= OTHER
/note= "2'-methoxyethyl nucleotides"
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/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 2 A; 3 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PARP-2 antisense inhibitor ISIS #126144.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                 Example 18; Page 91; 168pp; English
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/mod_base= OTHER
/note= "2' metho:
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les 16; Conservative
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*tag=
                                                                             WPI; 2001-602570/68.
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Best Local S
Matches 16
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Cowsert LM;

Monia BP,

WPI; 2001-090484/10.

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                                                                                                                                                                                                                                                                                                                                   The invention relates to antisense oligonuclectides targeted to human PARP nucleic acid and inhibiting expression of human PARP. PARP (Poly And Parenase) palsays an important role in chromatin decondensation, DNA replication, DNA repair, gene expression, malignant transformation, cellular differentiation and apoptosis. The antisense oligonucleotide inhibitors are useful for inhibiting the expression of PARP in human cells or tissues. They are also useful for treating a human with a disease associated with PARP especially hyperproliferative disorders (e.g. cancer), cellular injury resulting from oxidative stress, neurological (e.g parkinsonism, meningitis-associated intracranial complications and isolaemia), inflammatory and autoimmune disorders (e.g. arthritis) and diabetes. The present sequence is an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human hnRNP A1 phosphorothioate antisense oligonucleotide, SEQ ID NO:46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human hnRNP A1; heterogeneous nuclear ribonucleoprotein A1; heterogeneous nuclear ribonucleoprotein core protein A1; p40CRS; mRNA processing; transport; stabilisation; alternative splicing; donor splice site selection; telomere biogenesis; oncogenesis; apoptosis-associated protein; cancer; tumour formation; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                     Antisense compound useful for treating hyperproliferative, neurological, inflammatory and autoimmune disorders and diabetes inhibits human PARP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                          Example 16; Page 86; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1055 AGTCAATCCCAACAAGACA 1074
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                                                                                                                   02-MAR-2000; 2000US-00517467.
                                                                                  01-MAR-2001; 2001WO-US006572.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC92774 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 80.0
                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                      Popoff I, Cowsert LM;
                                                                                                                                                                                                                      WPI; 2001-602570/68.
                WO200164955-A1
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                                                   07-SEP-2001.
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Matches
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(ISIS-) ISIS PHARM INC

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cother beterogeneous nuclear ribonnucleoprotein A1 (hnRNP A1) gene, which to the heterogeneous nuclear ribonnucleoprotein A1 (hnRNP A1) gene, which to the heterogeneous nuclear ribonnucleotides were designed to target different regions of the human hnRNP A1 mRNA, and were analysed for their effect on hnRNP A1 mRNA levels by quantitative real-time PCR. hnRNP A1 (also Known as heterogeneous nuclear ribonnucleoprotein core protein A1 and p400RS) is thought to function in the stabilisation, transport and processing (including alternative splicing) of newly and shuttles continuously between the nucleus and the cytoplasm acting as a carrier protein for mRNAs. InRNP A1 also participates in telomere a carrier protein for mRNAs in hnRNP correlating with shortened a carrier protein for mRNAs a laso been classified as an apoptosis telomeres. In addition, hnRNP A1 has also been classified as an apoptosis control splicing events, particularly donor splice site selection, hnRNP control splicing events, particularly donor splice site selection, hnRNP control simplicated in the process of oncogenesis. The oligonnucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with hnRNP A1 expression, such as cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                  Novel antisense compound targeted to human hnRNP Al which specifically hybridizes with and inhibits the expression of human hnRNP Al, useful for modulating the expression of hnRNP Al in cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transport; membrane; cytostatic; virucide; vasotropic; dermatological; antipsoriatic; antiasthmatic; gene therapy; tumor cell; antisense; tumor therapy; drug; 8s.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 6 A; 12 C; 1 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-ICAM-1 oligonucleotide SEQ ID 53.
                                                                                                                                                                  Example 15; Col 41-42; 38pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               231 TGGTGGTGGTGGCGCCAGTG 250
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-203679/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-JUL-1999;
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Matches
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Secreted Frizzled-related protein; sFRP; chronic bronchitis; asthma; chronic obstructive pulmonary disease; COPD; antisense therapy; mouse; emphysema; reverse transcription PCR; RT-PCR primer; sfrp3 gene; s8.
                                                                                                                                                                                                                                                                                                                                                                     Mouse sfrp3 gene specific forward RT-PCR primer.
                                                                                                                                                                                                                                                                                        226 GAGAGTGGTGGTGGTGCCGG 245
                                                                                                                                                                                                                                                                                                   20 cacaceccaacrecreces 1
                                                                                                                                                                                                                                                                                                                                                                                                                                              28-FEB-2001; 2001WO-US006579.
                                                                                                                                                                                                                                                                                                                                  AAD17434 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                              AAD17434;
                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                      RESULT 1296
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This invention describes a novel conjugate (I) which consists of (A) a molecule to be transported and (B) at least one aryl residue of formula - (X - (Y - (Y + R))) in (II). Ar = group containing at least one aromatic ring; X = 0 or N (sic); X = 0 or N (sic); X = group containing at least one accomatic ring; X = 0 or N (sic); Y = 0, S or NHZ 2 (sic); R I = optionally substituted of the containing double and/or triple bonds); R 2 = optionally containing double and/or triple bonds; R 2 = optionally containing double and/or triple bonds; R 1 = integer of 1 or more. (A) is bonded to (B) directly or via a chemical group, provided that the chemical group; least one aryl residue (not restricted to containing a chemical group; R 2 = sective function at the invention also containing a reactive function at the position of the which (B'); by preparing (A') containing a reactive function at the position of the invention have cytostatic, proping (II) (optionally bonded via a chemical corporation and antiasthmatic activity and can be used for gene the invention have cytostatic, virucide, vasotropic, dermatological, antipsoriatic and antiasthmatic activity and can be used for gene therapy. Conjugation of (A) with (B) is useful for transporting (A) across biological membranes. The products of particularly tumor cells). Medicaments, diagnostic agents and test kits containing (I) are also claimed. Typically (I) are antisense or into enkaryotic or prokaryotic cells. (Specifically bacterial, yeast or mammalian cells, including human cells, containing (I) are also claimed. Typically (I) are antisense or into diagnostic agents and test kits containing (I) are also claimed. Typically (I) are antisense or into diagnostic agents and test kits containing (I) are also claimed. Typically (I) are antisense or into diagnostic agents and cellular uptake (C) (A), e.g. in tumor cells. (B) include fluorescein diactive to that obtained using other conjugates (I) are fluorescein diacetted (I) in the conjugates (I) are fluorescein diacetted (I) in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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New substituted aryl conjugates of parent molecules, especially oligonucleotides, having improved transmembrane and intracellular transport properties, useful as medicaments or diagnostic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
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                                                                                                                                                                                                                             Disclosure; Page 8; 28pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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therapy

The present sequence is mouse secreted Frizzled-related protein (sfrp3) gene specific reverse transcription PCR (RT-PCR) primer. The invention relates to a method for treating or preventing chronic obstructive pulmonary disease (COPD) such as emphysema, asthma and chronic bronchitis in a subject. The method involves administering to the subject, an agent effective to inhibit apoptosis by inhibiting the expression of a secreted Prizzled-related protein (sFRP) gene. It is also useful in antisense

Inhibition of apoptosis for the treatment or prevention of obstructive pulmonary disease comprises inhibiting expression of secreted Frizzled-

pulmonary disease comprises inhibiti related protein gene in lung cells.

(UYCO) UNIV COLUMBIA NEW YORK.

WPI; 2001-557764/62. D'armiento J, Imai

29-FEB-2000; 2000US-00514885.

Example 2; Page 35; 79pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is human secreted Frizzled-related protein 4 (sFRP4) gene specific reverse transcription PCR (RT-PCR) primer. The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Inhibition of apoptosis for the treatment or prevention of obstructive pulmonary disease comprises inhibiting expression of secreted Frizzled-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Secreted Frizzled-related protein; BFRP; chronic bronchitis; asthma; chronic obstructive pulmonary disease; COPD; antisense therapy; human; emphysema; reverse transcription PCR; RT-FCR primer; sFRP4 gene; ss.
                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                      0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.38+02;
iive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                  Sequence 20 BP; 8 A; 7 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human sFRP4 gene specific forward RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             related protein gene in lung cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 35; 79pp; English.
                                                                                                                                                                                                                                                                                                                                     890 ACATCATCAACATGCACAAC 909
                                                                                                                                                                                                                                                                                                                                                           1 ACATGACCAAGATGCCCAAC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYCO ) UNIV COLUMBIA NEW YORK.
                                                                                                                                                                                                                                                                                                                                                                                                                   AAD17410 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-FEB-2000; 2000US-00514885.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-FEB-2001; 2001WO-US006579.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                  16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            D'armiento J, Imai K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-557764/62.
                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200164717-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAD17410;
                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1297
                                                                                                                                                                                                                                                                                                                   Matches
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ВР

919

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The sequence represents PCR primer RP.2(rev) used in analysis of multiple tumour suppressor MTS1 and MTS2. The MTS genes, and expression products, are useful for treating, diagnosing or prognosing human cancer. In particular, the MTS gene is useful for diagnosing a predisposition to as a gene therapy for melanoma, leukaemia, astrocytoma, glioblastoma, hodghin's lymphoma, chronic lymphatic leukaemia (CLL), or cancers of the pancreas, breast, thyroid, ovary, uterus, testis, kidney, stomach or rectum. The gene may be used in both cancerous and pre
                                                                                                                                                                                                  ö
relates to a method for treating or preventing chronic obstructive pulmonary disease (COPD) such as emphysema, asthma and chronic bronchitis in a subject. The method involves administering to the subject, an agent effective to inhibit apoptosis by inhibiting the expression of a secreted Frizzled-related protein (SFRP) gene. It is also useful in antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New mouse multiple tumor suppressor gene, useful for diagnosing or prognosing human cancer or as gene therapy for treating cancer, particularly melanoma, leukemia, astrocytoma, lymphoma or cancers of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; multiple tumour suppressor; MTS1; MTS2; therapeutic; diagnostic; cancer; gene therapy; melanoma; leukaemia; astrocytoma; glioblastoma; lymphoma; glioma; Hodgkin's lymphoma; chronic lymphatic leukaemia; PCR primer; ss.
                                                                                                                                                                                                    Gaps
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0
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Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                            Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer RP.2 (rev) used in analysis of MTS1 and MTS2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                            Sequence 20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                       1289 TCCTGTCCAACGAGGAGTTC 1308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 7; Col 40; 80pp; English.
                                                                                                                                                                                                                                                                       rccreeccarceaecaerae 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95WO-US003316.
95US-00487033.
95US-00508735.
                                                                                                                                                                                                                                                                                                                                                                 AAS02589 standard; DNA; 20 BP
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Best Local Similarity 80.0%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ą;
                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jiang P, Kamb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-280859/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pancreas or breast.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancerous cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US6210949-B1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                         29-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stone S,
                                                                                                                                                                                                                                                                                                                                                                                                     AAS02589;
                                                                                             therapy
                                                                                                                                                                                                                                                                                                                              RESULT 1298
                                                                                                                                                                                                                                                                                                                                                 AAS02589,
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apoptosis; hyper-proliferating cell; cancer; tumour; eczema; cell-cycle progression regulator; genital warts; restenosis; skin cancer; psoriasis; scar tissue; intracellular-adhesion molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                 Inhibiting cancer cell proliferation by exposing cells to a composition of fusion proteins comprising VP22 polypeptides coupled to cell cycle progression regulators, and further exposing cells to cell death
                                                                                                                                             FITC; ICAM; oligonucleotide; ss; fluorescein isothiocyanate; VP22; BH3;
                                                                                                                                                                                                                                                /*tag= a
/note= "C is labeled with FITC"
                                                                                                                                                                                                                                                                                                                                                                             O'hare PFJ, Normand NM, Brewis ND, Phelan A;
                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Page 14; 23pp; English.
                                                                                                                            FITC-labeled ICAM oligonucleotide.
524
                -
           20 GAAGGCTICCIGGACACGCT
505 GAGGGCTACCTGGAGAAGCT
                                                                  AAS09545 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                    21-DEC-2000; 2000WO-GB004965.
                                                                                                                                                                                                                                                                                                                                       99GB-00030519
                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-418224/44
                                                                                                                                                                                                                                                                                                                                                           (PHOG-) PHOGEN LID.
                                                                                                                                                                                                                                                                              WO200147960-A1
                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                       24-DEC-1999;
                                                                                                                                                                                                                                        misc_feature
                                                                                                         24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                    stimulators
                                                                                                                                                                                                                                                                                                 05-JUL-2001
                                                                                                                                                                                                          Synthetic.
                                                                                       AAS09545;
                                                RESULT
                                                                    8 8
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The sequence represents an FITC (fluorescein isothiocyanate) labeled cligonucleotide complimentary to part of the mRNA encoding the intracellular-adhesion molecule ICAM. The oligonucleotide is included in a composition comprising a fusion protein of herpes virus VP22 protein 159-301 (laving the transport function) and a cell-cycle progression regulator (or its DNA) e.g. BH3 or apoptotic proteins. The composition is used to reduce the proliferation of cells. The method of making the VP22 containing compositions is used for reducing proliferation of hyperproliferating cells e.g., cancer cells, for manufacturing a medicament to reduce or treat cell proliferation e.g. cancer cell proliferation. The method is also used for reducing or treating cell proliferation. The method is also used for reducing or treating cell proliferation, in tumour cells present in tumour cell mass, non-malignant cells e.g., benigh tumour cells such as genital warts, smooth muscle cells present in restencist, proliferating cells e.g., skin cancer, psoriasis or eczema skin cells, or proliferating cells e.g., skin cancer, psoriasis or Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 9.3e+02; les 16; Conservative 0; Mismatches 4; Indels Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other; Query Match

226 GAGAGTGGTGGTGGCGG 245

Matches

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Gaps

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Gaps

12-JUN-2001 (first entry)

schultz621-3.rng

20 GAGAGGGGAAGTGGTGGGGG

g

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Convenient and cheap microplate fluorescent screening method for detecting gene anomaly in e.g. infectious diseases, congenital genetic diseases or cancers through gene diagnosis in community screening test
                                                                                                       HPV; genetic disease; gene anomaly; infectious disease; chlamydia; congenital genefic disease; cancer; human papilloma virus; k-ras; cystic fibrosis; mitochondrial cerebromyopathy; cervical cancer; colon cancer; PCR primer; ss.
                                                                                   PCR primer used to amplify a k-ras DNA sequence.
                                                                                                                                                                                                                                                                        (SAPP-) SAPPORO IMMUNO DIAGNOSTIC LAB
                                                                                                                                                                                                                                                                                             Yamaguchi A, Kikuchi K, Nakamura K;
                                                                                                                                                                                                                                                                                                                                                                                            Claim 7; Page 22; 26pp; Japanese.
                    ВР
                                                                                                                                                                                                                             09-FEB-2000; 2000WO-JP000693
                                                                                                                                                                                                                                                  09-FEB-2000; 2000WO-JP000693
                    AAH42979 Standard; DNA; 20
                                                              15-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                  WPI; 2001-497079/54.
                                                                                                                                                                                   WO200159124-A1.
                                                                                                                                                             Unidentified
                                                                                                                                                                                                         16-AUG-2001
                                         AAH42979;
RESULT 1300
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PCR primers AAH42977-80 were used to amplify k-ras DNA sequences. The primers are used in the method of the invention. The specification describes a method for screening genetic diseases. The method comprises using DNA simply extracted from a biological specimen such as scraped mucosal cells and tissue slide pieces fixed with formalin and embedded in paraffin, and amplifying a target region by polymerase chain reaction (PCR) for direct floursescence measurement of the additional doublestranded DNA intercalator. The method is used for detecting gene anomaly in e.g. infections diseases, congenital genetic diseases or cancers, including infection disease due to human papilloma virus and chlamydia genetic diseases like cystic fibrosis, mitochondrial cerebromyopathy, cancers of cervical cancer and colon cancer, through gene diagnosis in community screening test program
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.6; DB 1; Length 20;
llarity 80.0%; Pred. No. 9.3e+02;
Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 2 A; 1 C; 9 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1310 AGACATACAACTACCCCAAG 1329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
es 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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Matches
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555 CCTCAGCCGCCGCCTCCGTC 574
                                                                     20 cceccecceccecceccecci
                                                                                                                                              AAH41775 standard; DNA; 20
               16; Conservative
Local Similarity
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                                                                                                                                                                          AAH41775;
                                                                                                                  RESULT 1302
               Matches
                                            ò
                                                                     셤
                                                                                                                                                                                         20 ACACCTCCAACTACCACAAG 1
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Synthetic

AAF99116/c ID AAF99116 standard; DNA; 20 BP.

RESULT 1301

ò 요 AAF99116;

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The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulatory nucleic acid to a non-rodent subject is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine ritch (py-rich) or thymidine (1) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or or orthomyxoviridae), bacterial antigens (e.g. herpesviridae, retroviridae and/or or orthomyxoviridae), bacterial antigens (e.g. toxoplasma and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect in the contract of the packbone and to activate immune cells. Note: the
                                                                    Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infections disease, allergy, immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Base, string; tape; circular disc; ligand; immobilised; PCR primer; detection; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 0 A; 6 C; 14 G; 0 T; 0 U; 0 Other;
                                 Immunostimulatory nucleic acid #232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 101; Page 43; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               p38 gene PCR primer SEQ ID NO:22.
                                                                                                                                                                                                                                                                                                                            27-SEP-1999; 99US-0156135P
23-AUG-2000; 2000US-0227436P
                                                                                                                                                                                                                                                                           25-SEP-2000; 2000WO-US026383
                                                                                                                                                                                                                                                                                                                                                                                   (IOWA ) UNIV IOWA RES FOUND.
(COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Krieg AM, Schetter C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-273485/28
                                                                                                                                                                                                WO200122972-A2.
                                                                                                                                                                                                                                                                                                            25-SEP-1999;
27-SEP-1999;
                                                                                                                                                                                                                                       05-APR-2001.
                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH4177
ID AA
XXX
XXX
AAC
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The present invention describes bases in the shape of a string, tape or circular disc on the surface of which a plural number of different ligands are immobilised respectively in pre-determined domains. Also described are devices for detecting the binding between the ligands and receptors and methods for detecting the binding between the ligands are useful for detection in biochemical and diagnostic assays. The ligands are immobilised in line, so the user only needs to determine the presence or absence of receptor binding, without further processing. AAH41135 to AAH41115 represent primers which are used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antileukoprotease; ALP; secretory leukocyte proteinase; SLPI; human; cancer marker; ovarian tumour; ovarian-derived metastatic tumour; overexpression; low malignant potential tumour; ovarian carcinoma; serous carcinoma; mucinous carcinoma; endometrioid carcinoma; elear cell carcinoma; cancer; diagnosis; cytostatic;
                                                                                                                                                                                                                               String, tape or disk shaped bases with several different immobilized ligands including nucleic acids, sugars, peptides and proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human antileukoprotease (ALP) reverse PCR primer, SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 5 A; 5 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                    Example 1; Page 37; 56pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1236 ACACTICATCTICCGIAICT 1255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 AAAGTICATCTICGGCATCT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH23850 standard; DNA; 20 BP.
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                                                                           24-OCT-2000; 2000WO-JP007415.
                                                                                                        05-NOV-1999; 99JP-00315610.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   clear cell carcinoma; cancer
quantitative PCR primer; ss
                                                                                                                                     (TAKI ) TAKARA SHUZO CO LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-AUG-2001 (first entry)
                                                                                                                                                                       Kato I, Izu H, Asada K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYAR-) UNIV ARKANSAS
                                                                                                                                                                                                       WPI; 2001-343623/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200128500-A2.
             WO200135098-A1
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                                            17-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH23850;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH23850,
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The invention relates to methods for the diagnosis and treatment of ovarian tumours or ovarian-derived metastatic tumours in an individual. The diagnositic method involves measuring the level of antilenkoprotease (ALP) in a sample (e.g., a blood sample, tissue biopsy or ovarian secretion) from an individual. If the level of ALP exceeds the mean basal secretion from an individual is the level of ALP exceeds the mean basal constitutions, the individual is likely to have an ovarian or ovarian-constitution also known as secretory leukocyte protease inhibitor which specifically inhibits the activity of serratum corneum chymotryptic enzyme, and is also able to inhibit leukocyte elastise, cathepsin G, chymotrypsin and trypsin. It is significantly overexpressed in carcinomas methods of treating ovarian or ovarian-derived tumours, or preventing covarian tumour metastasis, via the administration of ALP. Methods of the ovarian and ovarian-derived metastatic tumours, particularly low malignant potential tumours or ovarian carcinomas such serous carcinoma. Consuminous carcinoma, endometriod are ovarian carcinoma, endometriod carcinoma, endometriod carcinoma and clear cell carcinoma. Sequences AAH23847-AAH23850 represent PCR primers used in quantitative consumed and cancerous ovarian tissue levels of ALP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                               Detecting tumor growth in an individual, particularly ovarian and ovarian derived metastatic tumors, comprises measuring antileukoprotease levels.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated unedited human excitatory amino acid 4 receptor polynucleotides and proteins, useful for screening potential therapeutic compounds and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer for nucleic acids encoding the human BAAS receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, excitatory amino acid 4 receptor, EAA 4 receptor;
central nervous system receptor, PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 7 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1109 CCCCTGACATCCTGCTTGGG 1128
                                                                                           Example 3; Page 10; 45pp; English.
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Best Local Similarity 80.0
Matches 16; Conservative
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WPI; 2001-290812/30
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0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ative 0; Mismatches 4; Indels

Underwood LJ, Shigemasa K;

Tanimoto H,

99US-0159972P

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human excitatory amino acid (EAA) 5 receptor. The synthesis this central nervous system (CNS) receptor in vivo is regulated by an editing mechanism. This editing results in the expression from a single human CNS receptor game of structurally distinct forms of the CNS receptor protein. The specification describes a human EAA4 receptor. The human excitatory EAA4 receptor polymuclacotide and the protein it encodes are useful for screening potential therapeutic compounds and selecting drug candidates that interact selectively with edited human central nervous system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse; multiple tumour suppressor; MTS; cytostatic; somatic mutation; germ line mutation; gene therapy; melanoma; leukaemia; astrocytoma; CLL; glioblastoma; lymphoma; glioma; Hodgkin's lymphoma; cancer; rectum; P16; pancreas; breact; thyroid; ovary; uterus; testis; kidney; stomach; mouse; P16beta; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to somatic and germ line mutations in the multiple tumour suppressor (MTS) gene in human cancer. The invention also relates to therapy of human cancer which have a mutation in the MTS gene,
                                                                                    primers AAC62056-60 were used to amplify nucleic acids encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel multiple tumor suppressor proteins useful for diagnosis and prognosis of human cancer and for screening drugs for cancer treatment.
 drug candidates that interact with edited human central nervous system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse P16beta cDNA amplifying P16-specific reverse PCR primer.
                                                                                                                                                                                                                                                                                                                  'Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 9.3e+02; les 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 4 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                          1211 CGGGCTCCACGGTGGAGGAA 1230
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                                                  Example 8; Col 21; 91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                               1 CTGGCTCCGAGGTGGTGGAA 20
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94US-00215086.
94US-00215087.
94US-00227369.
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95WO-US003316.
95US-00486047.
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18-MAR-1994;
18-MAR-1994;
14-APR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New regulatory sequences from the fascin gene, useful for providing dendritic cell-specific expression of e.g. antigens, e.g. for vaccination against tumors and infections.
including gene therapy, protein replacement therapy, and protein mimetics. The MTS sequences are useful for diagnosing predisposition to human cancer or for diagnosing and prognosing human cancers such as melanoma, leukaemia, astrocytoma, glioblastoma, lymphoma, glioma, Hodgkin's lymphoma, CLL and cancers of pancreas, breast, thyroid, ovary, uterus, testis, kidney, stomach and rectum. They are also used for screening drugs for cancer treatment. The present sequence is pl6-specific reverse PCR primer used for amplifying mouse Pl6beta cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antibacterial; antifungal; antiparasitic; anti-allergic; neurological; immunomdularory; apoptotic; expression regulator; vaccine; allergen; Creutzfeld-Jakob disease; Alzheimer's disease; gene therapy; autoimmune disease; transplant rejection; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fascin; regulatory sequence; human; dendritic cell; antiviral; tumor;
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Pred. No. 9.3e+02;
3; Mismatches 4; Indels
                                                                                                                                                                                    Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human fascin associated primer SEQ ID 55.
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02-MAR-2000; 2000DE-01010188.
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Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RESK) RESKB-KUNZ A.
(ROSS)) ROSS X.
(ROSS)) ROSS R.
(BROS/) BROS M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-451858/48.
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AAH48603/
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Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

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A new protein is described which comprises the 318 residue amino acid sequence corresponding to wild type retinol dehydrogenase (RDHS), but where amino acid 238 is not Gly, amino acid 73 is not Ser, or amino acid 33 is not Ile. This mutant RDHS can be used in the analysis of mutations in the gene encoding retinol dehydrogenase, in the diagnosis and treatment of ocular diseases associated with retinal degeneration such simple subjunctatus. Other disorders which may also be studied include fundus albipunctata albescens, albipunctate dystrophy and retinitis pigmentosa. A number of primer pairs (See GENSEEQ records AAA5443). A54448) were used to amplify the genomic RDHS DNA. Two primers (AAA54441, AAA54442) were used to amplify exon 3b of the RDHS gene. This primer corresponds to nucleotides 3151-3170 of the genomic DNA sequence (See GENESEQ record AAA5441).
in DC; for isolation and identification of cell factors and cis-elements from regulatory sequences that mediate DC-specific expression; to determine the degree of maturity of DC and to block transcription factors, by providing binding sites in DC. (A) provide DC-specific expression of nucleic acid under their control, allowing a more specific regulation of the immune response and eliminating the long and laborious purification of DC (since a complete leucocyte population may be transformed), including transformation in vitro. This sequence represents a primer associated with the human fascin gene described in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mutations in nucleic acid molecules encoding 11-cis retinol dehydrogenase correlated to ocular disorders, useful in diagnosis and treatment of diseases such as fundus albipunctatus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-cis retinol dehydrogenase; RDH5; eye; mutant; mutation; ocular disease; fundus albipunctatus; retinitis punctata albescens; albipunctate dystrophy; retinitis pigmentosa; human; primer; ss.
                                                                                                                                                                                                                                                            Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer for amplifying 11-cis retinol dehydrogenase (RDH5).
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                                                                                                                                                                                                                       Sequence 20 BP; 1 A; 10 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (LUDW-) LUDWIG INST CANCER RES.
(HARD ) HARVARD COLLEGE.
(MASS-) MASSACHUSETIS EYE & BAR INFIRMARY.
                                                                                                                                                                                                                                                                                                                                                    1631 CCAGCAGCCAGCGCTGGAG 1650
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                                                                                                                                                                                                                                                                                                                                                                                           20 ccadeadecedadecrecae 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to an antibody or its fragment that specifically binds to a human multiple tumour suppressor (WTS). The invention is useful for detecting differences in the absence of MTS peptides, to screen a tissue or to detect mutant MTS gene products. The antibodies will immunoprecipitate MTS proteins from solution as well as react with MTS protein on Western or immunoblots of polyacrylamide gels
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Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                   MTS; Multiple Tumour Suppressor; cancer; antibody; ss.
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                                                                                                                                                                                                                                       Primer used to amplify mouse beta cDNA.
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                                                      S6 TGTGACTGCTGAAACCCAGG 75
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94US-00215087.
94US-00227369.
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95WO-US003316
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18-MAR-1994;
14-APR-1994;
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17-MAR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes a method for detecting a mutation at a particular position in a target nucleic acid. The method comprises binding the target to a solid support, hybridizing a probe to the target, elongating the probe with nucleotide(8) resistant to exonuclease, digesting the probe with accounclease and detecting bound nucleic acid. The mutation is in position 'n' in a target nucleic acid and the 3' extremity of the probe hybridises to position 'n'. The method is used to detect gene mutations implicated in disease, particularly hereditary genetic diseases, especially sickle cell anemia alpha and beta cancer. The present sequence represents a probe which is used in the method of the invention to detect a mutation in codon 12 of K-ras
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, phosphofructokinase isozyme-2; iPFK-2; therapy; drug screening;
cancer; inflammation; cachexia; anti-tumour; phosphorothioate backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting mutation in target nucleic acid, useful for detecting hereditary genetic diseases, comprises using chip whose electrical or optical property changes relative to the presence of hybridized probe.
                                                                                                     DNA mutation; hereditary genetic disease; sickle cell anemia; probe;
thalassemia; cystic fibrosis; haemophilia; cancer; K-ras; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human iPFK-2 DNA specific phosphorothicate sense oligonuclectide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ouery Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                    Probe used to detect a mutation in codon 12 of K-ras.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 2 C; 10 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            229 AGTGGTGGTGGTGGCGCAG 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 5; Page 18; 36pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Acrestrecrestresascas 20
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                                                                                                                                                                                                                                                             01-MAR-2001; 2001WO-FR000604.
                                                                                                                                                                                                                                                                                              31-MAR-2000; 2000FR-00002614.
                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-557783/62.
                                                                                                                                                                                                                                                                                                                               (NUCL-) NUCLEICA
                                                                                                                                                                                        WO200164945-A2.
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Synthetic.
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                                 26-NOV-2001
                                                                                                                                                                                                                           37-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                Cailloux F;
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AAH78394;
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#XSSX#WWWEXPXPXBX

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The present invention relates to a method for screening a candidate therapeutic agent that inhibits kinase enzymatic activity of phosphofructokinase isozyme (IPRA-2). Phosphofructokinase catalyses the formation of fructose 2,6-biphosphate from fructose-6-phosphate. The method is used for identifying compounds that may be used to inhibit iPPR-2 activity, an enzyme that is over-expressed by cancerous cells. IPRK-2 is useful as diagnostic targets, drug screening targets and as antisense compounds that inhibit inflammation, cachexia and its translation in callular cytosol as an anti-tumour treatment. The present sequence is human phosphofructokinase isozyme (IPRK-2) DNA specific phosphorothicate sequence is sense oligonuclectide (S-IPRF-2) used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human iPFK-2 DNA specific phosphorothioate antisense oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, phosphofructokinase isozyme-2; iPPK-2; therapy; drug screening;
cancer; inflammation; cachexia; anti-tumour; phosphorothioate backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Screening for agents which inhibit the activity of the oncogenic phosphoglucomutase isozyme-2.
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                                                                                   /note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1679 CCAACTACATCTTCCCTGCT 1698
                                                                                                                                                                                                                                                                                                                                                                                                                                  Mitchell
1. .20
/*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                           (PICO-) PICOWER INST MEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                      97US-00961578.
                                                                                                                                                                                                                                                            98US-00183846.
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/*tag≕ a
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Chesney JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-424617/45.
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nes 16; Conserv
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modified_base
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Synthetic.
                                                                                                                                                                                                                                                         30-OCT-1998;
                                                                                                                                                                                                                                                                                                                      31-OCT-1997;
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                                                                                                                                             US6255046-B1
                                                                                                                                                                                                    03-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                     Bucala RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD11920;
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Example 1; Page 33; 152pp; English.

the protein.

schultz621-3.rng

03-JUL-2001

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The present invention relates to a method for screening a candidate therapeutic agent that inhibits kinase enzymatic activity of phosphofructokinase isozyme (IPFK-2). Phosphofructokinase catalyses the formation of fructose 2,6-biphosphate from fructose-6-phosphate. The method is used for identifying compounds that may be used to inhibit iPFK-2 activity, an enzyme that is over-expressed by cancerous cells. iPFK-2 is useful as diagnostic targets, drug screening targets and as antisense compounds that inhibit inflammation, cachexia and its translation in cellular cytosol as an anti-tumour treatment. The present sequence is human phosphofructokinase isozyme (iPFK-2) DNA specific phosphorothioate antisense oligonucleotide (AS-iPFK-2) used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Solute carrier family 6 neurotransmiter transporter; sectonin 4; SLC6A4; genotyping; allele specific oligonuclectide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated polynucleotide comprising a polymorphic variant for the solute carrier family 6 neurotransmitter transporter, serotonin member 4 gene for identifying drugs for treating disorders related to expression
                                                                                                                                                                                          Screening for agents which inhibit the activity of the oncogenic phosphoglucomutase isozyme-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                             Chesney JA, Mitchell RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (PICO-) PICOWER INST MEDICAL RES
                                                                                                                                                                                                                                           Example 4; Col 9; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
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                               98US-00183846.
                                                                97US-00961578.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JUL-1999; 99US-0146290P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF74084 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                              WPI; 2001-424617/45.
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                               30-OCT-1998;
                                                                31-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer #18.
                                                                                                                               Bucala RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF74084;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1312
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Gaps

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PCR primer used to amplify cDNA encoding a rat mu-subtype opioid receptor. The polynucleotide sequence is useful for producing a mu-type opioid receptor by standard recombinant techniques. The encoded protein is useful for producing monoclonal or polyclonal anti-receptor antibodies and to identify patterns of post-translational modifications and to elucidate associated G proteins. Mu receptor polynucleotides and solypeptides are useful in identifying other receptor subtypes, in screening for new opioid ligands and for understanding mechanisms of opioid action e.g., drug addiction
                                                                                                                                                                                                                                                                                             ö
                                                                   The present invention relates to a polymorphic variant of a reference sequence for the solute carrier family 6 neurotransmitter transporter, serotronin member 4 (SLC6A4) gene or a fragment of it or a sequence complementary to the first sequence. The invention is used in producing a recombinant organism that can be used to express SLC6A4 for protein structure analysis and binding studies. A composition comprising a general protein organization or structure analysis and binding studies. A composition comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated DNA encoding mu-subtype opioid receptor protein which is useful for identifying other receptor subtypes, screening for mu opioid ligands and for understanding mechanisms of opioid action.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer used to amplify cDNA encoding rat mu-subtype opiate receptor.
                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mu-subtype opioid receptor; G protein; opioid; drug addiction;
PCR primer; ss.
                                                                                                                                                                                                                                                       Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                217 GGCCTGGATGAGAGTGGTGG 236
                                                                                                                                                                                                                                                                                                                                                                1 Gecergeardadrecrered 20
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93US-00026140.
93US-00075447.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF85418 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-JUL-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-342395/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UHLG/) UHL G R.
(EPPL/) EPPLER C M.
(WANG/) WANG J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rattus rattus.
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26-FEB-1993;
11-JUN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF85418;
                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1313
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ö ö 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ative 0; Mismatches 4; Indels SQ Sequence 20 BP; 3 A; 5 C; 6 G; 6 T; 0 U; 0 Other; Ouery Match Best Local Similarity 80.0° Matches 16; Conservative

849 CCTGGACAAGGACCTGAAGC 868

20 ccreakakakarrakka 1

AAH49228 standard; DNA; 20 BP

AAH49228;

(first entry) 26-NOV-2001 Anti-ICAM oligonucleotide XXI.

Polyamide-oligonucleotide derivative, anticancer; antiproliferative, antiviral; hepatoropic; vasotropic; antisense inhibition; ribozyme; integrin; cell-cell adhesion; cancer; restenosis; stability; PNA; peptide nucleic acid; ss.

Synthetic.

EP1113021-A2

08-MAR-1995; 2001EP-00104012

14-MAR-1994; 94DE-04408528. 08-MAR-1995; 95EP-00103332.

(AVET) AVENTIS PHARMA DEUT GMBH.

Thlmann E, Breipohl G;

WPI; 2001-591267/67.

New DNA-peptide nucleic acid chimeras, useful e.g. as antisense agents for treating e.g. cancer, also as diagnostic probes and primers.

Disclosure; Page 24; 54pp; German.

This invention describes novel polyamide-oligonucleotide derivatives (I) and their physiologically acceptable salts of formula P((DNA-LI)_g(PNA-LI)_g(PNA-LI)_g(DNA-LI)_g(PNA-LI)_g(DNA-LI)_g(PNA-LI)_g(DNA-LI

more active than either DNA or PNA oligomers. When used as probes, (I) show different responses to base-pair mismatches in the DNA and PNA segments, allowing better discrimination between pathogenic and non-pathogenic conditions such as the transition from proto-oncogene no oncogene, also, when used as primers, with the PNA segment at the 5'-end, they produce amplicons resistant to 5'-exonuclease, allowing this enzyme to be used to eliminate RNA or DNA primers. The DNA component allows additional reactions not possible with PNA alone, e.g. 3'-tailing and (I) may be incorporated into a gene. AAH49208-AAH49264 represent oligonucleotides used to illustrate the method of the invention 888888888888888

Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;

ö 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ative 0; Mismatches 4; Indels Query Match Best Local Similarity 80.0° Matches 16; Conservative

ö

226 GAGAGTGGTGGTGGCGG 245 ò

20 GAGAGGGAAGTGGTGGGGG 1

RESULT 13 AAF87785/

AAF87785 standard; DNA; 20 BP

AAF87785;

(first entry) 11-JUL-2001 DNA 20-mer ASO (antisense DNA oligomer) SEQ ID NO:12.

Antisense DNA oligomer; ASO; identification; gene therapy; target; Nearest-Neighbour Thermal Stability Program; thermal melting temperature; phosphorothioate; disease treatment; DNA:RNA hybrid; ss.

Synthetic.

JS6183966-B1

06-FEB-2001.

99US-00235614. 22-JAN-1999;

94US-00320507. 97US-00808474. 03-MAR-1997;

(TEXA) UNIV TEXAS SYSTEM.

Gray DM, Clark CL;

WPI; 2001-280429/29

Identifying a nucleic acid having a sequence capable of targeting a gene of interest, for identifying nucleic acids for gene therapy, comprises using the Nearest-Neighbor Thermal Stability Program.

Example 1; Col 21-22; 43pp; English.

The present invention describes a method for the identification of a nucleic acid having a sequence capable of targeting a gene of interest compiratises: (a) a first database having a list of stability values for independent combinations of N(x); (b) a computing unit having a means for independent combinations N(x), data list, defining a nucleic acid sequence of interest to be targeted to provide a second database; and (c) sequence of interest. The method is useful for identifying a nucleic acid having a stability value of a nucleic acid sequence capable of targeting the gene of interest. The method is useful for identifying a nucleic acid having a sequence capable of targeting a gene of interest. The method may be used to obtain thermodynamic parameters for 20 combinations of nearest-neighbour base pairs of DNA:RNA hybrid sequences. The Nearest-Neighbour Thermal Stability Program can process data for use in

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calculating thermal melting temperatures for phosphorothioate DNA:RNA hybrids. The program can be readily extended to predict the most stable triplex-forming sequences, or antigene oligomers. The present sequence represents a DNA 20-mer ASO (antisense DNA oligomer) sequence which is used in the exemplification of the present invention
      8866666
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Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;

ö Gaps ô Ouery Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels

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AAF87788 standard; DNA; 20 BP AAF87788/

AAF87788;

(first entry)

11-JUL-2001

Human intracellular adhersion molecule 1 (ICAM-1) S-ASO SEQ ID NO:15

Antisense DNA oligomer; ASO; identification; gene therapy; target; Nearest-Neighbour Thermal Stability Program; thermal melting temperature; phosphorothicate; disease treatment; DNA:RNA hybrid; human; ICAM-1; intracellular adhersion molecule 1; ss.

Homo sapiens.

US6183966-B1,

06-FEB-2001,

99US-00235614. 22-JAN-1999;

97US-00808474. 94US-00320507 07-OCT-1994; 03-MAR-1997;

(TEXA) UNIV TEXAS SYSTEM.

Clark CL; Gray DM, WPI; 2001-280429/29.

Identifying a nucleic acid having a sequence capable of targeting a gene of interest, for identifying nucleic acids for gene therapy, comprises using the Nearest-Neighbor Thermal Stability Program.

Detecting collagen gene alteration, useful for diagnosing osteoporosis, multiple epiphyseal dysplasia, osteogenesis imperfecta, shortness of stature and low bone density in humans.

Claim 8; Fig 24; 617pp; English.

Example 1; Col 25-26; 43pp; English.

The present invention describes a method for the identification of a nucleic acid having a sequence capable of targeting a gene of interest comprises: (a) a first database having a list of stability values for independent combinations of N(x); (b) a computing unit having a means for inputting data comprising N(x), data list, defining a nucleic acid sequence of interest to be targeted to provide a second database; and (c) a program capable of processing the first and second database to N(x) comparison, and a stability value of a nucleic acid sequence capable of targeting the gene of interest. The method is useful for identifying a nucleic acid having a sequence capable of targeting a gene of interest. These mucleic acids are useful in gene therapy and disease treatment. The method may be used to obtain thermodynamic parameters for 20 combinations of nearest-neighbour base pairs of DNR:NR hybrid sequences. The Nearest-Neighbour Thermal Stability Program can process data for use in calculating thermal melting temperatures for phosphorothicate DNA:RNA hybrids. The program can be readily extended to predict the most stable program can be readily extended to predict the most stable triplex-forming sequences, or antigene oligomers. The present sequence represents an antisense DNA oligomer designated S-ASO targeted to the

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human intracellular adhersion molecule 1 (ICAM-1), which is used in an
                                                                                                                                                                                                                           Human; collagen; COLIA1; COLIA2; COL9A1; COL9A2; COL9A3; ss; osteoporosis; multiple epiphyseal dysplasia; osteogenesis imperfecta; shortness of stature; low bone density; gene therapy; PCR primer.
                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                     Koerkkoe J;
Paassilta P;
                                                                 ö
                                           Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                            Spotila LD, Deltas CD, Sereda L;
Larson A, Pack M, Colige A, Early J,
Annunen S, Pihlajamaa T, Vuoristo M,
                            Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                            Human COL9A2 PCR primer 1 for Exon 15.
          example from the present invention
                                                                                                                                                                                                                                                                                                                                                                        HEALTH SCI.
THOMAS.
                                                                                   226 GAGAGTGGTGGTGGCGG 245
                                                                                                     20 cacaccaacrecrecres 1
                                                                                                                                                     ВР.
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94US-00212322.
                                                                                                                                                                                                                                                                                                                           97US-00943731.
                                            0.8%;
                                                                                                                                                    AAS22310 standard; DNA; 20
                                                                                                                                                                                         (first entry)
                                                                 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                       (UYAL-) UNIV ALLECHENY
(UYJE-) UNIV JEFFERSON
(UYOU-) UNIV OULU.
                                             Query Match
Best Local Similarity
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Westerhausen
                                                                                                                                                                                                                                                                                                                                                                                                                                Ala-Kokko L,
                                                                                                                                                                                          24-OCT-2001
                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                           03-OCT-1997;
                                                                                                                                                                                                                                                                                                                                             03-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                       13-MAR-1994;
                                                                                                                                                                                                                                                                                                        24-JUL-2001.
                                                                                                                                                                      AAS22310;
                                                                                                                                  RESULT 1317
                                                                  Matches
                                                                                                                                             AAS223.
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The invention relates to Detecting a collagen gene alteration associated with a pathological condition in a human subject by obtaining from the subject a sample nucleic acid containing a portion of at least 15 consecutive nucleotides of the segment of the COLIA1 gene extending in the 5' to 3' direction from 78 nucleotides of intron 27 located adjacent exon 28 through the 3' end of intron 17, where the portion contains an intronic nucleotide and a first and second site, determining the sequence of the portion contains an intronic nucleotide and a first and second site, determining the sequence of the portion with the corresponding consensus sequence of the COLIA1 gene where a difference between the sequence of the portion and the consensus sequence indicates the presence of the collagen alteration in the subject. The method is used for detecting abnormalities in a COLI or COL9 gene is useful for determining whether a subject is afflicted with pathological conditions associated with an altered collagen gene such as osteoporosis, multiple epithyseal dysplasia, osteogenesis imperfecta, shortness of stature and consumently in a collagen gene is also useful for designing a therapeutic nucleotide or gene therapy agent which can be administered to the subject to correct or alleviate the

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ABL01636;
                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Use of aggregates comprising VP22 protein/polypeptide with the transport function of VP22 and oligonucleotides/polynucleotides with disaggregating agent, useful for treating or preventing cell proliferation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to the use of aggregates comprising VP22 (viral protein 22) protein (or a polypeptide with the transport function of VP22), and oligomuclectides or polymucleotides with a disaggregating agent e.g. Aluminium phthalocyanine (AT) (simultaneously or sequentially) to treat target cells by delivering molecules to the cells and/or preventing cell proliferation and/or killing cells. Also included are a method of treating target cells to deliver molecules to the cells and/or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'note= "C is covalently linked to a fluorescein moiety"
                                                                                                                                                                                                                                                                                                                                                         VP22; viral protein 22; ss; cytostatic; antipsoriatic; dermatological; disasgregating agent; Aluminium phthalocyanine; cell proliferation; apoptosis; psoriasis; eczema; skin cancer; restenosis; scarring; Intracellular-adhesion molecule; ICAM.
abnormality. The method is useful for detecting mutations in both the coding and non-coding sequences of any of the COLI or COL9 genes. Therefore the method can be used to detect collagen gene alterations which affect either the primary sequence of a collagen protein chain, splicing of the mRNA encoding such chains or regulation of expression the genes encoding such chains. The present sequence is a PCR primer which amplifies a nucleic acid from a collagen gene of the invention
                                                                                                                                                  Gapa
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                                                                                                                       0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ve 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                   Intracellular-adhesion molecule, ICAM, oligonucleotide.
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A-tag= A-THER
/note= "Phosphorothioate backbone"
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                                                                                                 Seguence 20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                        1632 CAGCAGGCAGCGGCTGGAGG 1651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 17; 31pp; English
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/label= OTHER
                                                                                                                                                                                                                                                             ВР
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                                                                                                                                    80.08
                                                                                                                                                                                                                                                            ABK12803 standard; DNA; 20
                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                    Best Local Similarity 80.0
Matches 16; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
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                                                                                                                                                                                                                                                                                    ABK12803;
                                                                                                                           Query Match
                                                                                                                                                                                                                                      RESULT 1318
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prevent their proliferation and/or kill them comporising: (a) exposing the cells to the aggregate composition cited above; and (b) exposing the disaggregation of the aggregate cited above; which can promote disaggregation of the aggregate composition in cells, where steps (a) and the aggregate composition and the disaggregating agent, as combined preparation for administration of these components, either sequentially or together, a pharmaceutical comprising the aggregating agent composition and cort disaggregating agent, in combination with a pharmaceutical excipient and a cell preparation obtainable by treating the target cells in vitro as cited in the method above. The aggregate composition and treating diseases or target cells, and/or preventing cell proliferation cort creating diseases or target cells, and/or preventing cell proliferation and creating therapy, or as a medicament for delivering medicaments for useful in therapy, particularly for manufacturing medicaments for use in therapy, or as a medicament for delivering molecules to cells to prevent cell proliferation or kill cells. In particular, these may be used for treating psoriasis, eczema, skin cancer, restenosis and scarring. The composition code present sequence is an oligomuche encoding an Intracellular-adhesion contents.
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/note="This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide nucleic acid, PNA, cytostatic, virucide, dermatological, antiasthmatic, overexpression, viral infection, vitiligo, antisense, pigmentation disorder, asthma, polyamide backbone, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ICAM-1 targeted antisense peptide nucleic acid SEQ ID NO: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.6; DB 1; Length 20; 0.0%; Pred. No. 9.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      226 GAGAGTGGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AVET ) AVENTIS PHARMA DEUT GMBH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            80.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-MAR-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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The invention relates to detecting (MI) infection by human herpes virus (HW) by performing informatics analysis of gene sequences from different HHV types or strains (e.g. HHV)-HHVS) to identify target segment (TS), selecting probe and primers capable of directing amplification, amplifying TS, interpolating HHV number by comparing number of amplification cycles (NAC) for detecting TS to NAC to detect known quantity of TS. Also included are cloning a segment of genomic viral DNA
                                                                                                                                           The present invention relates to peptide nucleic acid (PNA) derivatives having at the C-, and optionally N-, terminus one or more phosphoryl groups, at least one of which contains one or more deprotonisable groups, preferably hydroxy or mercapto. These PNAs are useful in the treatment of tumours or any disease associated with (over) expression of particular genes, including viral infections, vitiligo or other pigmentation disorders, and asthma. The present sequence is a peptide nucleic acid described in the exemplification of the invention
                                New peptide nucleic acid derivatives, useful e.g. for tumor treatment and diagnosis, contain terminal, deprotonizable phosphoryl groups for e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting infection of human herpes virus type or strain by informatic analysis of gene sequence using probe and primers capable of directing amplification of target sequence and interpolating the virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human herpes virus infection; ss; real time PCR; primer; HHV1; HHV2; HHV3; HHV4; HHV5; HHV6; HHV7; HHV8; latent membrane protein-1; LMP-1; nuclear protein EBNA2; intermediate early protein; IE; glycoprotein B;
                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HHV4a nuclear protein EBNA2 forward real time PCR primer.
                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                             Disclosure; Page 22; 93pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                     226 GAGAGTGGTGGTGGTGCGG 245
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(first entry)
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hes 16; Conservative
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                                                                        improved solubility.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human herpesvirus 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-463369/49.
WPI; 2002-075055/10.
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26-AUG-2002
                                                      diagnosis,
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critical the ledentified TS (MZ), a polymucleotide (1) molecule having any one crown the identified TS (MZ), a polymucleotide an HHV1 thymidine kinase protein from a drug-resistant HHV2 thymidine kinase protein from a drug-resistant HHV2. thymidine kinase protein from a drug-resistant HHV2, thymidine kinase protein from a drug-resistant HHV2, thymidine kinase protein, HHV4 latent membrane protein. I, an HHV4 con an HHV4b latent membrane protein. I, an HHV4 con a drug-resistant HHV2, thymidine kinase protein. I, an HHV4 con a drug-resistant HHV2, HHV4b latent membrane protein. I, an HHV6 conclear protein. I, an HHV6 conclear protein. I, an HHV6 conclear protein EBNAZ, HHV4b latent membrane protein. I, an HHV6 conclear protein. HHV6 glycoprotein Bonaz, and an HHV6 intermediate early protein, an HHV6 dlycoprotein Bonaz, HHV6 intermediate early protein, an HHV6 dlycoprotein Bonaz, and an HHV6 intermediate early protein, an HHV7 glycoprotein Bonaz, enclear protein glycoprotein (i.e. the target sequences), and a fluorescence quecher group covalently attached to the probe, and a fluorescent reporter group covalently attached to the probe, (MX) is useful for cloning (MZ) a segment of artached to the probe, (MX) is useful for cloning (MZ) a segment of particular type or a strain of HHV in a sample from an individual containing the flooredures of mallyse the effectiveness of pharmaceuticals by meanuring the ability of anti-viral agents to mediate HHV propagation. (MI) allows accurate and procedures, infection by one strain of a specific type of HHV type of anti-viral agents of the problems endamining specific act of anti-viral agents, purification by HHV that cannot be detected by conventional PCR approaches. In addition to determining specific act of anti-viral agents, purification by more strain of a specific order with a crommer problems endamine prover the present of sequence last he hHV4 an uclear protein EBNAZ forward real time PCR primer. Con anti-viral agents of sequence last is bunavailable. The present is seque
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the identified TS (M2), a polynucleotide (I) molecule having any one
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es 16; Conservative
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Gaps ö

AAD41528 standard; DNA; 20 30-OCT-2002 (first entry) AAD41528; AAD41528/

Marker, vitamin D analogue, antiproliferative, cancer, osteodystrophy; multiple sclerosis, osteoporosis, osteomalacia; hyperparathyrotdism; genoprotective, epidermal wound; chemoprotective; DNA repair mechanism; cytostatic; psoriasis; neuroprotective; Vulnerary; RT-PCR; primer; ss.

Collagenase 1 gene specific reverse RT-PCR primer.

28-NOV-2001, 2001WO-CA001689 WO200244403-A2. Unidentified 06-JUN-2002.

29-NOV-2000; 2000US-0253746P 02-MAY-2001; 2001US-0287729P (UYMC-) UNIV MCGILL

WPI; 2002-537458/57 White JH;

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The invention relates to a marker for testing analogues of vitamin D expected to be effective in reducing aberrant activity of vitamin D responsive cell, comprises at least one gene pertinent to the action of vitamin D for testing analogues and determining analogues capable of regulating the gene, and is indicative of a chemopreventive or chemotherapeutic again. The invention is useful for testing analogues of vitamin D expected to be effective in reducing aberrant activity of vitamin D-responsive cell or for testing analogues of vitamin D suspected to have antipizoliferative activity. The invention is useful for reducing aberrant activity of vitamin D-responsive cell, and for treating a disorder characterised by an aberrant activity of vitamin D-responsive cell, where the disorder is selected from cancer, psoriasis, multiple cell, where the disorder is selected from cancer, psoriasis, multiple cell, where the disorder is selected from cancer, psoriasis, multiple cell, where the disorder is selected from cancer, psoriasis, multiple cell, where the disorder is selected from cancer, psoriasis, multiple cell, where the disorder is useful for identifying regulated the invention is useful for protecting against in vivo DNA damage to the skin of a mammal, or for reducing or preventing DNA damage to the skin of a mammal, preferably human. The invention is useful as a marker for the activity of DNA repair mechanisms. The invention is useful as a marker for the activity of DNA repair minibiting an enzyme which metabolises 1.25-dihydroxyvitamin D3 The invention is useful for testing compounds susceptible of inhibiting an enzyme which metabolises 1.25-dihydroxyvitamin D3 The invention is useful for treating epidermal wounds. The present sequence invention is useful for treating epidermal wounds. The present sequence is collagenase 1 gene specific RT-PCR primer
                                          Novel marker for testing analogs of vitamin D expected to be effective in reducing aberrant activity of vitamin D-responsive cell, comprises gene pertinent to action of vitamin D for testing the analogs.
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80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
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(NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY
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                                                                                                                                                                                                                                                                                    Example 2; Page 48; 89pp; English
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Matches 16; Conservative
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EXEXE EXECUTE 
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New isolated human or mouse targeting peptide useful for targeted

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                                                                                                                                            The invention relates to the screening of antitumour agents by using the interaction between ARP protein and HK33 (Housekeeping 33) protein.

Nuclear transport of ARF protein is inhibited by the expression of HK33 gene, and thus p53-dependent transcription is suppressed. In immortalised cells, moreover, the expression of HK33 gene is significantly elevated. The invention provides a method of screening an antitumour agent by using the interaction between ARP protein and HK33 protein. It also provides a method for utilisation of HK33 protein and agene encoding it in the examination of tumour related disease. The current sequence represents a ARF/HK33 protein related primer
delivery of therapeutic agents, for inhibiting angiogenesis, tumor growth or pregnancy, and for inducing apoptosis or weight loss.
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80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
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                                                                                                   Example 6; Page 76; 81pp; Japanese.
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The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule (II) encoding a natural dominant negative regulator of caspase 8, FilP-c, where (I) specifically hybridises with and inhibits expression of the protein, or specifically hybridises with at least an 8-nucleobase portion of an active site on (II). (I) has antiinflammachory and anti-tumour activities. (I) is an inhibitor of FilP-c expression, a modulator of apoptosis and can be used in antisense gene therapy. (I) is useful for inhibiting the expression of FilP-c in cells or tissues, and for treating an animal having a disease or condition associated with FilP-c. (I) is also useful for modulating apoptosis in a cell, where a caspase such as caspase 8, caspase 3 or caspase 7 is activated, and the FilP-c is the long form of FilP-c. (I) is also useful for diagnostics, therapeutics, prophylaxis, as research reagents and kits, for distinguishing functions of various members of biological pathway, and in antisense gene therapy. (I) is also useful brophylactically, e.g., to prevent or delay infection, inflammation or tumour formation. The present sequence represents mouse FilP-c inhibiting chimeric phosphorothylactically in four formation. The present sequence represents mouse FilP-c inhibiting command of the prophylactic phonomial or command the command of the prophylactic phonomial or command the command of the prophylactic phonomial or command the command or command the command or command the command or command the command or wings, which is used in an example from the present invention

Sequence 20 BP; 2 A; 6 C; 4 G; 8 T; 0 U; 0 Other;

Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels 661 TACAAAGGCAAAAGCAAGCT 680 20 TACACAGGCAGAGGT 1 à

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Gaps

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ABQ74294 standard; DNA; 20 BP. ABQ74294; RESULT 1324

14-OCT-2002 (first entry)

Human leukocyte antigen DQB1 locus PCR primer DQB1-ex2F.

Human leukocyte antigen; DQB1; DQA1; aspermia; examination; detection; PCR primer, ss.

Homo sapiens.

JP2002153300-A.

28-MAY-2002.

24-NOV-2000; 2000JP-00358486.

24-NOV-2000; 2000JP-00358486.

(INOK/) INOKO H.

WPI; 2002-552748/59.

Examination of aspermia comprising investigating an allele with correlation to aspermia if it is detected in the HLA-DQA1 locus.

Example 2; Page 4; 7pp; Japanese.

The present invention describes a method for the examination of aspermia in which, if an allele showing correlation to aspermia is detected in the human leukocyte antigen (HLA)-DQA1 locus, it is investigated. Also described is a method for the examination of aspermia in which one of the following (a) to (e) is investigated: (a) if the base sequence of the DNA corresponding to codon 64 of HLA-DQA1 gene is AGA; (b) if the base sequence of the DNA corresponding to codon 66 of HLA-DQA1 gene is ATG; (c) if the base sequence of the DNA corresponding to codon 69 of HLA-DQA1 gene is GTG; (d) if the base sequence of the DNA corresponding to codon 69 of HLA-DQA1 gene is GTG; (d) if the base sequence of the DNA corresponding to codon 69 of HLA-DQA1 gene is GTG; (d) if the base sequence of the DNA corresponding to codon 69 of HLA-DQA1 gene is GTG; (d) if the base sequence of the DNA corresponding to codon 69 of HLA-DQA1 gene is GTG; (d) if the base sequence of the DNA

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corresponding to codon 71 of HLA-DQA1 gene is GTG. The method is useful for the examination of aspermia. The present sequence represents a PCR primer for the HLA-DQB1 locus, which is used in an example from the present invention
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                                                                                                                                                  0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                           Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
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Best Local Similarity 80.0
Matches 16; Conservative
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Human, mouse, SAC1, carbohydrate, sweetener; ethanol, alcoholism, ss;
obesity, diabetes, transgenic embryo, body tissue, body fluid, pancreas,
blood; tongue, PCR primer, anorectic, antidiabetic, gene therapy;
                                                                                                                                                                                                          Human SAC1 gene-specific oligonucleotide PCR primer #45.
                                               AAS97894 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-APR-2000; 2000US-0200794P.
28-JUL-2000; 2000US-0221419P.
10-NOV-2000; 2000US-0247443P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-APR-2001; 2001WO-US013387
                                                                                                                                                                                                                                                                                                                                          protein replacement therapy.
                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200183749-A2.
                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                     12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-NOV-2001.
                                                                                                      AAS97894;
  1325
RESULT 13
AAS97894
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Novel isolated polypeptide comprising variant form of mouse or human SAC1 polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.

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De Jong PJ,

Chatterjee A, Tordoff MG;

Bachmanov AA, Beauchamp GK, Ohmen JD, Reed DR, Ross D,

WPI; 2002-075162/10.

(WARN) WARNER LAMBERT CO. (MONE-) MONELL CHEM SENSES CENT.

Claim 14; Page 91; 239pp; English.

The invention relates to an isolated polypeptide, comprising a variant for form is associated with altered preference for carbohydrates, other sweeteners or ethanol.

The polypeptide and its associated DNA sequence can be produced by recombinant techniques and its useful for preventing obesity, disbetes or alcoholism associated with SACI expression. The sequences are useful in screening for drugs and sweeteners. Recombinant call lines and transgenic correcting for drugs and sweeteners. Recombinant call lines and transgenic morpor may be used in screening for and identifying agents that induce or repress function of SACI. Predisposition to diabetes, obesity or alcoholism can be ascertained by testing any fluid or tissue of a human such as blood, pancreas or tongue) for sequence variations of the SACI gene. A sequence variation of the SACI locus may indicate a predisposition to diabetes, obesity and/or alcoholism and may provide a predisposition to diabetes, obesity and/or alcoholism and may provide a sample by contacting the DNA with a probe to form a hybridiaation complex which is then detected. The sequences represent cDNA encoding human and

(first entry)

23-APR-2002

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      highest expression gene group consisting of 100 genes which show the highest expression among the genes expressed in human maturation/ activation DC. Also described are: (1) a protein expressed by the above human maturation/activation DC expression gene; (2) an antibody against the protein; and (3) an antagonist against the expression of each gene belonging to the above gene group. The gene group is useful for the treatment and the diagnosis of various human diseases related to human DC. ABL42927 to ABL42956 represent PCR primers for human maturation/ activation DC expression genes, which are used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                      Human; maturation/activation dendritic cell expression gene; maturation; activation; dendritic cell; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a human maturation/activation dendritic
                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                      Maturation/activation dendritic cell expression gene PCR primer #328.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
mouse SAC1 polypeptides and PCR primers specific for the SCA1 genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human maturation/activation dendritic cell expression gene group.
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                                                                   Length 20;
                                                                                                 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 11 C; 2 G; 4 T; 0 U; 0 Other;
                              Sequence 20 BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                tch 0.8%; Score 13.6; DB 1; al Similarity 80.0%; Pred. No. 9.3e+02; 16; Conservative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 39; 41pp; Japanese.
                                                                                                                                  851 TGGACAAGGACCTGAAGCAG 870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                481 CTACCAGCTGACATCGGGCT 500
                                                                                                                                                         1 TGGAGTACGACCTGAAGCTG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  creccascreacercaeer 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-MAY-2000; 2000JP-00150562
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Best Local Similarity 80.0
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-127070/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP2001327293-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens
                                                                                                                                                                                                                                                                                                                      12-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                    ABL42954;
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                                                                Query Match
                                                                                 Local
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                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modulating the expression of human glioma-associated oncogene-1. The antisense compounds, particularly antisense oligomucleotides, target and inhibit the expression of human glioma-associated oncogene-1. The antisense compounds are useful for inhibiting the expression of human glioma-associated oncogene-1. The glioma-associated oncogene-1 in human cells or tissues and for treating an animal, particularly a human suspected of having or being prone to a disease or condition associated with expression of glioma-associated oncogene-1. The compounds are useful for diagnostics, therapeutics and as research reagent, e.g. prophylactically to prevent or delay infection, inflammation or tumour formation. The antisense compounds are safely and effectively administered to humans. ABK10509-ABK30586 represent the antisense oligomucleotides of the invention which comprise a phosphorothioate backbone
                                                     Human glioma-associated oncogene-1 antisense oligonucleotide ISIS 124842.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Angiogeneeis inhibitor; ss; angiogeneeis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint;
                                                                                        Human, glioma-associated oncogene-1 associated disease, infection,
inflammation; tumour formation; cytostatic; antiinflammatory; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel antisense compounds targeted to nucleic acids encoding glioma-associated oncogene-1, for modulating the gene expression and treating diseases associated with expression of the oncogene in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    present invention relates to antisense compounds and methods for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 5 A; 4 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anglogenesis inhibitory oligonucleotide #243.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             340 GACTIGAAGAIGGGGICTGA 359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Col 44; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABS77759 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%;
                                                                                                                                                                                                                                                                                      08-SEP-2000; 2000US-00657042.
                                                                                                                                                                                                                                                                                                                             08-SEP-2000; 2000US-00657042,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Beet Local Similarity 80.0%;
...hes 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                         Bennett CF, Wyatt J;
                                                                                                                                   phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-138363/18.
                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                            US6329203-B1
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administering at least one antianglogenic nucleic acid molecule. Also nucleic acid sa kit comprising a first container housing the antianglogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted anglogenesis. The method is tuesful for inhibiting anglogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, conneal graft rejection, neovascular glaucoma, retrolencal fibroplasia, neovascularisation, telangiectasia, haemophiliac joints, anglofibroma, wound gramulation, intestinal adhesions, atherosclerosis, scleroderma and acid of the invention
                                                                                                                                                                                                                                                                                                         Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to inhibiting angiogenesis in a subject,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 6 C; 14 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunostimulatory nucleic acid SEQ ID NO: 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     555 CCTCAGCCGCCGCCTCCGTC 574
                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 23; 276pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 CCGCCGCCGCCGCCGCC 1
                                                                                                                                                                                                            (COLE-) COLEY PHARM GROUP INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABL39008 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-JUN-2000; 2000US-0213346P.
                                                                                                                                             14-DEC-2001; 2001WO-US048458.
                                                                                                                                                                            14-DEC-2000; 2000US-025534P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-JUN-2001; 2001WO-US020154.
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                                                                             WO200253141-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-APR-2002
                                                                                                                                                                                                                                            Bratzler RL;
                                                                                                              11-JUL-2002
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                                                Synthetic
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The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, olon and rectum cancer, connective tissue cancer, oesophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, kodgkin's lymphoma, nalanoma, myeloma, oral cavityr cancer, oversidan cancer, pancreatic cancer, prostate cancer, rabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "OTHER= Phosphorothioate internucleotide linkages,
bases 1-5 and 16-20 are 2'-methoxyethoxy (2'-MOE) bases.
All cytidine bases are 5-methylcytidines"
                                                                                          comprises
                                                                                                               administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, mouse, Protein Phosphatase 2 catalytic subunit alpha; diabetes; cancer; infection; inflammation; tumour formation; cytostatic; antidiabetic; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human/mouse Protein Phosphatase 2 antisense oligonucleotide #7.
                                                                                                such as basal cell carcinoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 0 A; 6 C; 14 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                Disclosure; Page 199; 312pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              555 CCTCAGCCGCCGCCTCCGTC 574
                                                                                                                                                                                                                                                                                                                                                                                                                              exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ccecceccecceccecce 1
                                                                                                Treating or preventing cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              멾.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%;
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(IOWA ) UNIV IOWA RES FOUND.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16; Conservative
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                                  Ö
                                 Hartmann
                                                                WPI; 2002-154611/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                               developing cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                 Weiner G,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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peptide nucleic acid derivatives, useful e.g. for treating tumors and mosis, have N-terminal phosphoryl residue for improving e.g.
             WPI; 2002-657604/70.
                                                                                                                                                                                                                                                                                                     WPI; 2002-089643/12.
                                                                                                                                                                                                                                                                                                                   diagnosis, have N-te
solubility in water.
                                                                                                                                                                                                                                                WO200179249-A2
                                                                                                                                                                                                                                  Unidentified
                                                                                                                                                                                               16-APR-2002
                                                                                                                                                                                                                                                         25-OCT-2001
    Monia BP,
                                                                                                                           Query Match
Best Local Si
Matches 16,
                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                     ABA97491;
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The present invention relates to peptide nucleic acid (PNA) derivatives. These can be used in the treatment of cancer, viral infections, vitiligo or other pigmentation disorders, and asthma. The present sequence is an oligonucleotide fragment of a PNA described in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to the cDNA and protein sequences of novel proteins HYPIND1 or FCHL1 and to sequence variations within these genes that have been shown to be associated with lipid disorders.

Oligonucleotide probes that hybridise to the CDNA sequence are useful for analysing the expression of FCHL1 by detecting the expression of the mRNA transcript in the sample. A host cell transformed with the CDNA of the invention is useful for producing the protein by recombinant means. Pharmaceutical compositions based on the sequences of the invention are useful for treating or preventing a lipid disorder associated with expression of FCHL1 such as familial combined hyperlipidaemia, coronary extery disease, atherogenic lipoprotein phenotype attacks of coronary extery disease, atherogenic lipoprotein phenotype familia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse; primer; antilipaemic; cardiant; hypotensive; anorectic; HYPLIP1; FCHL1; lipid disorder; familial combined hyperlipidaemia; coronary artery disease; atherogenic lipoprotein phenotype; cancer; hyperapobetalipoproteinaemia; hypertriglyceridaemia; obesity; ss; familial dyslipidaemic hypertension; syndrome X; insulin resistance; hypercholesterolaemia; chromosome 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel HYPLIPI and FCHL1 genes and their sequence variations associated with lipid disorder and cancer, useful for prognosis, diagnosis and treatment of lipid disorders.
                                                                                                                                                                                                                 Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       De Jong P,
                                                                                                                                                                Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse HYPLIP1 locus specific primer 273L17S #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chatterjee A,
, Wu C;
                                                                                                                                                                                                                                                                                                             226 GAGAGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 11; Page 76; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                        20 chadaccanacracracacca 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-SEP-2000; 2000US-0231322P
                                                                                                                                                                                                                                        80.08;
                                                                                                                                                                                                                   0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK68325 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Castellani LW,
oss D, Tafuri
                                                                                                                                                                                                                                                               16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-339808/37.
                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ross D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200220847-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bodnar JS,
                                                                                                                     invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ohmen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK68325;
                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                    The present invention relates to antisense oligomucleotides and methods for modulating the expression of human or mouse Protein Phosphatase 2 catalytic subunit alpha. The antisense oligomucleotides are useful for inhibiting the expression of Protein Phosphatase 2 catalytic subunit alpha and for treating diseases or conditions associated with aberrant expression of Protein Phosphatase 2 catalytic subunit alpha and for treating diseases or conditions associated with aberrant diseases include diabetes and cancer. The antisense oligomucleotides are also useful for diagnostics, therapeutics, and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. They are also useful as research reagents for distinguishing between functions of various members of a biological pathway. ABS65400-ABS65477 represent human or mouse Protein Phosphatase 2 catalytic subunit alpha antisense oligonucleotides which comprise a phosphorothioate backbone
                                                                                                        New antisense oligonucleotides targeted to nucleic acid encoding Protein Phospharase 2 catalytic subunit alpha, useful in treating diseases associated with the aberrant expression of Protein Phosphatase 2 catalytic subunit alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide nucleic acid, PNA, polyamide backbone; phosphoryl radical; cytostatic; virucide; dermatological; antiasthmatic; cancer; antisense; viral infection; vitiligo; pigmentation disorder; asthma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 6 A; 4 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                           Claim 3; Page 94; 153pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            51 AGCAGTGTGACTGCTGAAAC 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGCAGTGTAACTGTTTCAAC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AVET ) AVENTIS PHARMA DEUT GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Uhlmann E, Breipohl G, Will DW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              491/c
ABA97491 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16; Conservative
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               Wyatt JR;
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hyperapobetalipoproteinaemia, hypertriglyceridaemia, familial dysliplidaemic hypertenation, syndrome X, obesity, insulin resistance and hypercholesterolaemia. The CDNA sequence is useful in the diagnosis or prognosis of predisposition to lipid disorders and cancers, and also to

Disclosure; Page 87; 96pp; German

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Gaps

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The invention relates to a compound of 8-50 nucleobases in length targeted to a nucleic acid encoding protein phosphatase 1B (PTP1B), where the compound specifically hybridises with and inhibits the expression of PTP1B (e.g. an antisense oligonucleotide). Also included are (1) a compound of 8-50 nucleobases in length which specifically hybridises with an a nucleobase portion of an active site on a nucleic acid encoding PTP1B; (2) inhibiting the expression of PTP1B in cells or tissues comprising contacting the cells or tissues with the compound; treating an animal having or usupected of having a disease or condition associated with PTP1B comprising administering the compound; (4) decreasing blood sugar levels in an animal comprising administering the compound; (5) preventing or delaying the onset of a disease or condition associated with PTP1B in an animal comprising administering the compound; (6) preventing or delaying the onset of an increase in blood glucose levels
identify a molecule which enhances or decreases the HYPLIP1 or FCHL1 activity. The present sequence represents an oligomucleotide primer specific for the mouse HYPLIP1 locus of the invention. The mouse HYPLIP1 locus is situated on chromosome 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Compound for inhibiting the expression of protein phosphatase 1B (PTP) and for treating diabetes, cancer, or obesity, comprises an antisense oligonucleotide targeted to nucleic acid encoding PTP1B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense; protein phosphatase 1B; PTP1B; se; probe; human; type 2 diabbetes; obseity; ovarian cancer; chronic myeloid leukaemia; hyperproliferative disease; antidiabetic; anorectic; cytostatic; blood glucose; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Freier SM, Monia BP, Butler MM, Mckay R;
                                                                                                                              / Match
0.8%; Score 13.6; DB 1; Length 20;
Local Similarity 80.0%; Pred. No. 9.3e+02;
tes 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PTP1B antisense oligonucleotide ISIS 142051.
                                                                                              Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                            16 GGATGGACAGGAATGCAGAG 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 27; 133pp; English.
                                                                                                                                                                                                                                    20 GGATGGAGAGGCATCCTGAG 1
                                                                                                                                                                                                                                                                                                                                        ABK85293 standard; DNA; 20 BP,
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31-JUL-2000; 2000US-00629644.
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                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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FREIER S M.
MONIA B P.
BUTLER M M.
MCKAY R.
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                                                                                                                                                                                                                                                                                                                                                                              ABK85293;
                                                                                                                                    Query Match
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(FREI/)
(MONI/)
(BUTL/)
(MCKA/)
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ABK85293/c
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in an animal comprising administering the compound. The compound is used to inhibit the expression of PTPLB in cells or tissues, to treat or prevent or delay the onset of a disease or condition associated with PTPLB, such as type 2 diabetes, obesity, cancer (especially ovarian cancer, chronic myeloid leukaemia and hyperproliferative diseases in an animal having or ususpected of having the disease or condition, and for decreasing blood sugar levels or preventing or delaying the onset of an increase in blood glucose levels in an animal. The compound is also used kits. The present sequence is an antisense compound of the invention targetting human PTPLB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel antisense compound targeted to nucleic acid encoding Fas, Fas
ligand or Fas associated protein-1 is useful for inhibiting expression of
Fas, Fas ligand, or Fap-1 in cells or tissues, and for treating
hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to an antisense compound encoding Fas, Fas ligand, or Fas associated protein-1 (Fap-1). The inhibition of Fas mediated signalling is thought to be immunosuppressive, antiinflammatory, heparotropic, cytostatic and vasotropic. Antisense oligonucleotides were designed to target human Fas. Oligonucleotides were synthesised as chimaric oligonucleotides and are useful for treating an animal having an autoimmune or inflammatory disease e.g., hepatitis, cancer, a condition associated with apoptosis, allograft rejection, or ischemia reperfusion injury. Optionally, the above mentioned conditions are prevented by contacting the allograft with the antisense oligonucleotide. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, immunosuppressive; antiinflammatory; hepatotropic; cytostatic; vasotropic; hepatitis; cancer; allograft rejection; ds; Fas.
                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                              0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
iive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human FasL chimeric phosphorothioate oligonucleotide #14.
                                                                                                                                                                                                                            Sequence 20 BP; 3 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zhang H;
                                                                                                                                                                                                                                                                                                                                          727 GAGGGGCACCCTGCACCGC 746
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18-SEP-2000; 2000US-00665615.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                            Query Match 0.8°
Best Local Similarity 80.0°
Matches 16; Conservative
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(MARC/) MARCUSSON E G
(WYAT/) WYATT J.
(ZHAN/) ZHANG H.
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ABQ79631 standard; DNA; 20 BP.
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                                                                     25-NOV-2002
                                                                                                                                                                                                            02-JUL-2002.
                                                                                                                                                                                                                                                                                                       Bucala RJ,
                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Si
Matches 16
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                                               ABQ79631;
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   RESULT 1336
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to antisense oligonucleotides of at least 10 bases complementary to inducible human phosphofructokinase-2 (iPFK-2) cDNA. The antisense oligonucleotides can be included in anticancer or antiinflammatcry pharmaceutical compositions along with an oligonucleotide carrier. An iPFK-2 antisonist such as an enzymatic inhibitor, anti-iPFK-2 antibody, or iPFK-2 antisense molecule can be administered for treating inflammatory disease or rapidly-growing cancers. The present sequence represents an iPFK-2-specific sense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel antisense oligonucleotides useful for treating inflammatory diseases or cancers, comprises complementary sequence of inducible human
oligonucleotides are used in diagnostics, therapeutics, prophylaxis and as research reagents and in kits. The oligonucleotides are also useful for research purposes. The present nucleotide sequence is related to
                                                                                                                                                                                                                                                                                iPFK-2-specific oligonucleotide S-iPFK-2 (A) (sense, position 35-55).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                     Human, phosphofructokinase-2; iPFK-2; antisense therapy; anticancer;
antinflammatory; cytostatic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.6; DB 1; Length 20; llarity 80.0%; Pred. No. 9.3e+02; Conservative 0; Mismatches 4; Indels
                                                                               Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02;
                                                                                                       4; Indels
                                                         Sequence 20 BP; 3 A; 10 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mitchell RA;
                                                                                                                            1659 CACCCCTCACAGGGCAGCCC 1678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1679 CCAACTACATCTTCCCTGCT 1698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (PICO-) PICOWER INST MEDICAL RES.
                                                                                                                                            CCCTCTTCACATGGCAGCC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 4; Col 8; 28pp; English.
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                                                                                                                                                                                                           ABQ79630 standard; DNA; 20 BP
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                                                                              0.8%;
                                                                                                                                                                                                                                                        (first entry)
                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bucala RJ, Chesney J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              phosphofructokinase-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-641574/69
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                                                                                          Local Similarity
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16;
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                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                     16;
                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                   ABQ79630;
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                                                                               Query Match
                                                                                                                                                                                     RESULT 1335
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Matches
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Matches
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The invention relates to antisense oligonucleotides of at least 10 bases complementary to inducible human phosphofructokinase-2 (1PFK-2) cDNA. The antisense oligonucleotides can be included in anticancer or antiinflammatory pharmaceutical compositions along with an oligonucleotide carrier. An iPFK-2 antagonist such as an enzymatic inhibitor, anti-iPFK-2 antibody, or iPFK-2 antisense molecule can be administered for treating inflammatory disease or rapidly-growing cancers. The present sequence represents an iPFK-2-specific antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense oligonucleotides useful for treating inflammatory diseases or cancers, comprises complementary sequence of inducible human phosphofructokinase-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                           phosphofructokinase-2; iPFK-2; antisense therapy; anticancer;
                                               iPFK-2-specific oligo AS-iPFK-2 (A) (antisense, position 35-55).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ô
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Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human chromosome 1p36-35 PCR primer SEQ ID NO:1374.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chesney J, Mitchell RA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1679 CCAACTACATCTTCCCTGCT 1698
                                                                                                                                                                                                                                                                                                                                                                                                                                         (PICO-) PICOWER INST MEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 CCAACGGCATCTTCGCGGCT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Col 25; 28pp; English.
                                                                                                                        antiinflammatory; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%;
                                                                                                                                                                                                                                                                                                                                           97US-00961578,
                                                                                                                                                                                                                                                                                                                                                                                             97US-00961578.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABL44330 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-APR-2002 (first entry)
(first entry)
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les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-641574/69.
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WPI; 2002-144136/19. (GENO-) GENOTEX YG.

10-MAR-2000; 2000JP-00066716. 12-MAR-2001; 2001JP-00068285.

20-NOV-2001.

(RIKA) RIKAGAKU KENKYUSHO. (GENO-) GENOTEX YG.

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ABT13935;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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11D ABT1

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                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates i(b) a primer described based on the chromsome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal overseponding to the marker is detected from the resultant plates containing the clones having said marker sequence; (d) the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to plates; (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed respectively in each wells of longitudinal discrimination Nos. are mixed respectively in each wells of longitudinal resultant cultures are amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL4257 to ABL45322 represent represent FOR primers for human chromosome 124221, which are represent for human chromosome 124221, which are represent corporation in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human chromosome 1p36-35 PCR primer SEQ ID NO:602.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 9 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Page 32; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      397 GAGGIGCAGICTCCAGIGAG 416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 GAGGIGAATGCTGCAGTGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL43558 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-MAR-2000; 2000JP-00066716.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 80.0%
                                                                                                                                                                                                                                                                                                                           Arraying genome clones.
                                                                                                                                                                                                                                                                       VPI; 2002-144136/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP2001321190-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1338
ABL43558/c
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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates independ for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. gucceed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed respectively in each wells of longitudinal discrimination Nos are mixed respectively in each wells of longitudinal creation Nos are mixed respectively in each wells of longitudinal discrimination Nos are mixed respectively in each wells of longitudinal creation the amplified products; (h) the clones in the multiwell plates are specified from the detected fromes are cultured and the created from the detected soult; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABI45357 to ABI45322 represent represent POR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, antisense gene therapy, phosphorothioate backbone, antisense oligonucleotide, helicase-moi gene; inflammation; ss; helicase-moi-associated condition; infection; tumour formation; 2-MOE nucleotide; 2'-methoxyethyl nucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  / Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 9.3e+02; hes 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human helicase-moi inhibiting oligonucleotide #60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 7 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                     Claim 4; Page 16; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       519 GAAGCTGACCCTCAATAGCC 538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 GAAGATGACGCTGAAGAGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABT13935 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-FEB-2003 (first entry)
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                                                                                                              Arraying genome clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-749291/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ward DT, Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6444466-B1.
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The invention comprises antisense oligonucleotides which are targeted to the coding region of the human helicase-moi gene. The antisense oligonucleotides of the invention are useful for inhibiting the expression of human helicase-moi in cells or tissues, and for treating a helicase-moi-associated condition. The antisense oligonucleotides of the invention may also be used to delay infection, inflammation and tumour formation. The present DNA sequence represents a human helicase-moi gene antisense oligonucleotide of the invention. NOTE: The present DNA sequence has a phosphorothioace backbone, bases 1-5 and 16-20 are 2'-methoxyethyl (2'-MOE) nucleotides
and for treating inflammation, specifically hybridizes to a specific region in nucleic acid molecule encoding the human helicase-moi.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 3 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                   Example 15; Col 45-46; 52pp; English.
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; Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels

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1380 GGCCGACCTCCTCACCAAGC 1399

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20 GGACTACCTCATAACCAAGC 1

RESULT 1340

AAI67702 standard; DNA; 20 AAI 67702

BP.

AA167702;

(first entry) 27-FEB-2002 SHH patched receptor (Ptc) cDNA amplifying forward primer.

Cell culturing, embryonic stem; ES; central nervous system; Ptc; Shh; dopaminergic; cholinergic; serotonergic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; tranquilizer; vulnerary; neuroleptic; cerebroprotective; cell therapy; gene therapy; CNS; PCR primer; ss.

Homo sapiens

40200183715-A2

08-NOV-2001.

01-MAY-2001; 2001WO-US014051

01-MAY-2000; 2000US-0201005P

US GOVERNMENT. LEE S. LUMELSKY N.

(USGO) US GOVERNMENT (LEES/) LEE S. (LUME/) LUMELSKY N. (STUD/) STUDER L. (MCKA/) MCKAY R D G.

Studer L, Mckay RDG; Lumelsky N, see S,

WPI; 2002-049345/06.

Culturing cells such as neuronal cells for use in treating neurological disorders, comprises generating embryoid bodies from undifferentiated embryonic stem cells, selecting precursor cells, expanding and differentiating them.

Example 10; Page 40; 66pp; English

The invention provides a method of culturing cells. The method involves expanding a culture of undifferentiated embryonic stem (ES) cells, generating embryoid bodies (EB), culturing the bodies to select for central nervous system (CNS) precursor cells (PC), culturing PC in an

expansion medium comprising a neurologic factor, and differentiating and culturing the expanded PC to form a culture of differentiated neuronal cells useful for culturing undifferentiated SS cells to form differentiated sscalls to form differentiated neuronal cells which are useful for treating a neurological disorder, especially Parkinson's disease in a patient. A gene product such as tyrosine hydroxylase, nerve growth factor (NDF), brain derived neurotrophic factor (BDNF), brGF, glial derived growth factor (GDNF) NT-3, and NT-4/5 can be introduced into a brain of a subject. The method is useful for culturing dopaminergic, cholinergic and service neuronal cells. The differentiated neuronal cells are useful for treating neurological disorders such as Euntington's disease, alreading epilepsy, familial dysauconomia as well as injury or trauma to the nervous system such as neurotoxic injury or disorders of mood and such as stroke and CNS disorders insuling from aging. Assays are useful for developing drugs capable of regulating the survival, profileration or comprise 20-40% dopaminergic neurons and 1-3* astroytes are useful for studying the mechanism of neuronal cells and to screen for an anging. Assays and selection neuronal cells and to screen pristing 50* 85* neurons which companies of neuronal cells and to screen for agonist of studying the mechanism of neuronal cells and release, particularly for serotonin and dopamine, neuronal cell survival, and the electrophysiochemical properties of differentiated neuronal cells.

CS Sequences AAI67692-721 represent genes, used for examining the companient of ES cells

Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;

Gape .; 0 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ive 0; Mismatches 4; Indels st Local Similarity 80.0 stches 16; Conservative Query Match Matches

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RESULT 1341 ABL46178/c

ABL46178 standard; DNA; 20 BP

ABL46178;

26-APR-2002 (first entry)

Human ICAM-1 antisense oligonucleotide ISIS 1939 SEQ ID NO:145.

Nucleic acid accessible hybridisation site; detection; hybridisation; characterisation; identification; nucleic acid structure; diagnosis; PCR primer; probe; ss.

Homo sapiens. Synthetic.

WO200198537-A2.

15-JUN-2001; 2001WO-US019401 27-DEC-2001.

17-JUN-2000; 2000US-0212308P.

(THIR-) THIRD WAVE TECHNOLOGIES INC.

Vener IT; Lyamichev V, Allawi H, Dong F, Neri BP,

WPI; 2002-049698/06.

Identifying oligonucleotides hybridizing to nucleic acids containing secondary structure, useful in clinical diagnosis, comprises identifying primers that interact with the target to form an extension product under

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23-MAY-2001; 2001WO-IB001153.
                                                                       retardation.
                                  ABK24601;
                              RESULT 13
ABK24601/
                                ð
                          g
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The present invention describes a method for identifying oligonucleotides with desired hybridisation properties to nucleic acid targets containing accondany structure. The method comprises amplifying a target nucleic acid having at least one accessible and one inaccessible site. Primers that form an extension product are identified as the oligonucleotides which can interact with the folded target nucleic acid. Oligonucleotides from the present invention can be used in novel detection methods for clinical diagnostic purposes, including the detection and identification of pathogenic organisms (e.g. HIV). The method allows the ability to rapidly analyse nucleic acid structures. ABL46034 to ABL46367 represent enemptication of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                     Example 17; Page 382; 409pp; English.
amplification conditions.
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tch 0.8%; Score 13.6; DB 1; Length 20; al Similarity 80.0%; Pred. No. 9.3e+02; 16; Conservative 0; Mismatches 4; Indela 226 GAGAGTGGTGGTGGCGG 245 Best Local Similarity Matches 16; Conserv

20 cadadedeaadredredede 1

ABK24601 standard; DNA; 20 BP

EIF2AK3 gene sequencing primer #17. (first entry) 09-APR-2002

Human; EIP2AK3; antidiabetic; osteopathic; antiarthritic; hepatotropic; nephrotropic; notropic; diabetes; Wolcott-Rallison syndrome; WRS; osteoporosis; arthritis; hepatic dysfunction; nephropathy; renal dysfunction; mental retardation; primer; ss; eukaryotic initiation factor 2 alpha kinase 3.

Homo sapiens.

WO200190371-A1.

29-NOV-2001.

23-MAY-2000; 2000EP-00401436. 02-OCT-2000; 2000EP-00402707.

(INRM) INSERM INST NAT SANTE & RECH MEDICALE. (NAGE-) CENT NAT GENOTYPAGE.

Nicolino M; Julier C, Delepine M,

WPI; 2002-122021/16

New mutated eukaryotic initiation factor 2 alpha kinase 3 genes and polypeptides in patients with Wolcott-Rallison syndrome, useful for preventing or treating e.g. diabetes, osteoporosis, arthritis or mental

Example 4; Page 31; 93pp; English.

The invention relates to an isolated variant of a mammal genomic sequence of the gene coding for the translation initiation factor 2 alpha kinase 3 (EIPIZAK3). The EIPIZAK3 nucleic acid variant is useful for the production of a recombinant or synthetic polypeptide, and for screening compounds capable of modulating EIPIZAK3. The nucleic acid is also useful for

screening or diagnosing the diseases cited below. The nucleic acid of may be used as sense or anti-sense oligonucleotide. The nucleic acid may also be used as primer or a probe, for detecting and/or amplifying a nucleic acid sequence. The compound is useful as a medicament, particularly for a roid sequence. The compound is useful as a medicament, particularly for preventing and/or treating diabetes and/or pathology related to WRS, e.g. type I diabetes, type 2 diabetes, the others forms of diabetes, c.g. cype 2 diabetes, the others forms of diabetes, c.g. creal dysfunction, or mental retardation, nephropathies or other can dysfunction, or mental retardation. The cell the mammal or the ETP2AX3 protein, and the direct or indirect interactions between the ETP2AX3 protein, and chemical or biochemical compounds, which may be a supersection of the tell for screening chemical or biochemical compounds capable of interacting directly or indirectly with the ETP2AX3 protein, and/or interacting the expression or the activity of the ETP2AX3 capable of modulating the expression or the activity of the ETP2AX3 coding sequences and the protein. ABX24521-ABX24624 represent human ETP2AX3 coding sequences and the primers of the invention ö Gaps ö 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels Sequence 20 BP; 6 A; 3 C; 5 G; 6 T; 0 U; 0 Other; 16; Conservative Query Match Best Local Similarity Matches

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Gaps ö

ABT06761 standard, DNA; 20 RESULT 1343 ABT06761,

ABT06761;

BP.

(first entry) 07-NOV-2002 Nucleic acid detection and discrimination related oligo SEQ ID No 104.

Hybridising, quantification; detection; synthesis; amplification; oligonucleotide; ds.

Unidentified.

WO200257479-A2.

25-JUL-2002.

27-DEC-2001; 2001WO-US050460.

27-DEC-2000; 2000US-00748146. 23-OCT-2001; 2001US-0330468P.

(INVI-) INVITROGEN CORP.

Darfler M; Nazarenko I, Rashtchian A, Solus J, Pires RM, Gebeyehu G, Astatke M;

WPI; 2002-627370/67.

Composition comprising nucleic acid molecules and a oligonucleotide capable of hybridizing with a portion of nucleic acid, and comprises modified nucleotide at or near the 3'-terminal nucleotide.

Example 29; Page 158; 307pp; English.

The invention relates to a composition comprising one or more nucleic acid molecules and at least one oligonucleotide, where at least a portion of the oligonucleotide is capable of hybridising with at least a portion of the nucleic acid molecule and where the oligonucleotide comprises a modified nucleotide at or near the 3'-terminal nucleotide. The various analogue oligonucleotides are useful for quantification or detection of one or more target nucleic acid molecules in a sample during nucleic acid

Page 637

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a composition comprising one or more nucleic acid molecules and at least one oligomucleotide, where at least a portion of the oligomucleotide is capable of hybridising with at least a portion of the nucleic acid molecule and where the oligomucleotide comprises a modified nucleotide at or near the 3'-terminal nucleotide. The various analogue oligomucleotides are useful for quantification or detection of one or more target nucleic acid molecules in a sample during nucleic acid synthesis or amplification. The analogues are also useful for determining the presence or absence of one or more particular nucleotides at a specific position or positions in a target nucleic acid molecule. The analogue oligomucleotides can also be useful for synthesising or amplifying one or more nucleic acid molecules, by mixing one or more nucleic acid molecules, by mixing one or more nucleic acid molecules or more nucleic acid cubating the mixture to synthesise or amplify one or more nucleic acid
synthesis or amplification. The analogues are also useful for determining the presence or absence of new or more particular nucleotides at a specific position or positions in a target nucleic acid molecule. The analogue oligonucleotides can also be useful for synthesising or manaplating one or more nucleic acid molecules, by maxing one or more nucleic acid molecules, by maxing one or more nucleic acid molecules, or an analogue oligonucleotides, and incubating the mixture to synthesise or amplify one or more nucleic acid molecules opposed and particular to all or a portion of the templates or targets. This polynucleotide sequence represents a nucleic acid detection and discrimination related oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Composition comprising nucleic acid molecules and a oligonucleotide capable of hybridizing with a portion of nucleic acid, and comprises modified nucleotide at or near the 3'-terminal nucleotide.
                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid detection and discrimination related oligo SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hybridising, quantification, detection, synthesis, amplification, oligonucleotide, ds.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Solus J, Pires RM, Darfler M;
                                                                                                                                                                                                           0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                              Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 29; Fig 36; 307pp; English
                                                                                                                                                                                                                                                                          948 CTACTGCCACGGCAGAAGG 967
                                                                                                                                                                                                                                                                                                         20 CTACAGCCACCATGAGAAGG 1
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23-OCT-2001; 2001US-0330468P.
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Gebeyehu G, Astatke M;
                                                                                                                                                                                                                                                                                                                                                                        ABT06751/c
ID ABT06751 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                             Query Match
Best Local Similarity 80.0
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (INVI-) INVITROGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-627370/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200257479-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JUL-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                      ABT06751;
                                                                                                                                                                                                                                                                                                                                                         RESULT 1344
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The invention relates to a composition comprising one or more nucleic acid molecules and at least one oligonucleotide, where at least a portion of the oligonucleotide is capable of hybridising with at least a portion of the nucleic acid molecule and where the oligonucleotide comprises a modified nucleotide at or near the 3'-terminal nucleotide. The various analogue oligonucleotides are useful for quantification or detection of comprises or more target nucleic acid molecules in a sample during nucleic acid comprises or expected acid molecules are also useful for determining the presence or absence of one or more particular nucleotides at a specific position or positions in a target nucleic acid molecule. The analogue oligonucleotides can also be useful for synthesising or analogue oligonucleotides can also be useful for synthesising or nucleic acid templates or targets with the analogue oligonucleotides, and incubating the mixture to synthesise or amplify one or more nucleic acid molecules complementary to all or a portion of the templates or targets. This polynucleotide sequence represents a nucleic acid detection and discrimination related oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                  ö
molecules complementary to all or a portion of the templates or targets. This polynucleotide sequence represents a nucleic acid detection and discrimination related oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid detection and discrimination related oligo SEQ ID No 103.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Composition comprising nucleic acid molecules and a oligonucleotide capable of hybridizing with a portion of nucleic acid, and comprises modified nucleotide at or near the 3'-terminal nucleotide.
                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                  4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                   0.8%; Score 13.6; DB 1;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pires RM,
                                                                                                                                                     Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Solus J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 29; Page 158; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                         948 CTACTGCCACCGGCAGAAGG 967
                                                                                                                                                                                                                                                                                                                                                                                                                                               CTACAGCCACCATGAGAAGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    멾
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-DEC-2000; 2000US-00748146.
23-OCT-2001; 2001US-0330468P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nazarenko I, Rashtchian A,
Gebeyehu G, Astatke M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABT06760 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                  16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-627370/67.
                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200257479-A2.
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0.8%; Score 13.6; DB 1; Length 20;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention comprises antisense oligonucleotides designed to inhibit expression of Syntaxin 4 interacting protein. The antisense oligonucleotides of the invention are useful for inhibiting the expression of Syntaxin 4 interacting protein in cells or tissues. The antisense oligonucleotides are also useful for treating an animal having a disease or condition associated with Syntaxin 4 interacting protein (e.g. diabetes, obesity or a skeletal muscle disorder). The antisense oligonucleotides can also be used to prevent or delay infection, franchamenton and tumour formation. The present DNA sequence represents a human Syntaxin 4 interacting protein antisense oligonucleotide. NoTE: The present percent sequence contains a phosphorothioate backbone and 2'-0-
                                                                                                                                                                                                                                                                                                                        Human, antisense gene therapy, Syntaxin 4 interacting protein; ss; antisense oligomucleotide; diabetes; obesity; skeletal muscle disorder; inflammation; tumour formation; phosphorothioate backbone; 2'-0-methoxyethyl wing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel antisense compound that hybridizes and inhibits nucleic acid molecule encoding Syntaxin 4 interacting protein, useful for treating diabetes, obesity and skeletal muscle disorder.
                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                       Human syntaxin 4 interacting protein antisense oligonucleotide 76.
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80.0%; Pred. No. 9.3e+02;
ative 0; Mismatches 4; Indels
                   4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seguence 20 BP; 11 A; 2 C; 1 G; 6 T; 0 U; 0 Other;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4
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                                                    948 CIACTGCCACCGGCAGAAGG 967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page 84; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 darricaaaaaararaacra 20
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                                                                                20 CTACAGCCACCATGAGAAGG 1
                                                                                                                                                                             ABQ62337 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ31505 standard; DNA; 20
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Best Local Similarity 80.0°
Matches 16; Conservative
                   16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monia BP, Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-404952/43.
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200224864-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                    16-AUG-2002
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                                                                                                                                                                                                                 ABQ62337;
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                                                                                                                                            RESULT 1346
                   Matches
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ID ABZ3
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ID ABQ
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The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprishing modifying one allele by insertion or replacement by a cassette having an expressible selectable marker and modifying other allele by recombination, of a promoter replacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are modified. The diploid fungal cells having both alleles modified are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that contributes to the virulence and/or pathogenicity of a fungus, a gene that contributes to the resistance of a fungus to an antifungal agent that inhibits the growth of a diploid fungus or and for identifying a compound which modulates the activity of a gene product, preferably enrymatic activity, carbon activity of a gene product, preferably enrymatic activity, carbon activity. The method is useful for identifying a compound which modulates the activity. The method is useful for identifying a compound having the ability to inhibit growth or proliferation of C. albicans cells and for treating infection by C. albicans. The present esquence is that of a present in the method of the invention. Note: The sequence is that of a present in the method of the invention. Note: The sequence is that of a present in the method of the invention. Note: The sequence is that of the method is the present entering the present parameter of a per present in the method of the invention. Note: The sequence is that of the present is not represented in the present esquence is that of the present in the present of the present parameter of a per present of the present parameter of a per present of the present parameter 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         on sequence information supplied to Derwent by the Buropean Patent Office
                                                                                                                                       Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 13.6; DB 1; Length 20; 10.0%; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Indels
                                                                                           Candida albicans GRACE strain PCR primer SEQ ID NO 5724.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ohlsen KL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 0 A; 0 C; 11 G; 9 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bussey H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       231 TGGTGGTGGTGCGCAGTG 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Boone C,
                                                                                                                                                                                                                                                                                                                                                                                                                                    29-DEC-2000; 2000US-0259128P.
20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
                                                                                                                                                                                                                                                                                                                                                                                          26-DEC-2001; 2001WO-US049486.
                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ELIT-) ELITRA PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 16; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jiang B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-566694/60.
                                                                                                                                                                                                                                             Candida albicans.
                                                                                                                                                                                                                                                                                            WO200253728-A2.
                                               30-JAN-2003
                                                                                                                                                                                                                                                                                                                                         11-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1348
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ID ABA
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ABA99824 standard; DNA; 20

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This invention describes a novel murine calpain protease 12 (capn12). The calpain protease of the invention, related proteins and mucleic acid that encodes it, are useful for treatment (including gene therapy) of diseases associated with insufficient expression of the calpain protease. The specific immunoglobulins (Ig) useful for diagnosis. Also the polymucleotide encoding capn12 is useful, e.g. as primers and probes, for diagnosis of diseases, or predisposition to them, and for recombinant production of capn12. This sequence represents the murine calpain 12, capn12 exon 19 splice donor site described in the disclosure of the
                                                                                                     Calpain protease; murine; gene therapy; screening; diagnosis; capn12; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                New calpain protein 12 with cysteine protease activity, useful for treating specific deficiency disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ouery Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 3 A; 6 C; 4 G; 7 T; 0 U; 0 Other;
                                                                       Murine capn12 exon 19 splice donor site.
                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 2c; 36pp; German.
                                                                                                                                                                                                                                                                                                                                              30-JUN-2000; 2000DE-01031932.
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/*tag= b
/number= 19
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/*tag= a
/number= 19
                                          (first entry)
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                                          11-JUN-2002
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              ABA99824;
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Polynucleotide and polypeptide of human NEDD-1 useful for diagnosing, treating or preventing a disorder associated with decreased or increased expression or activity of the polypeptide.

30-JAN-2001, 2001WO-US000663. 30-JAN-2001, 2001WO-US000663. 30-JAN-2001, 2001WO-US000664. 30-JAN-2001, 2001WO-US000665. 30-JAN-2001, 2001WO-US000666. 30-JAN-2001, 2001WO-US000667. 30-JAN-2001, 2001WO-US000669.

30-JAN-2001; 2001WO-US000670 01-JUN-2001; 2001US-00872462

(AEOM-) AEOMICA INT.

WPI; 2002-426011/45 Gu Y, Corrigan A;

26-SEP-2001; 2001WO-US030287

WO200226818-A2.

schultz621-3.rng

04-APR-2002

Example 2; Page 94; 190pp; English

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This invention relates to an isolated polynucleotide encoding human NEDD-1, which is cytostatic in its action. The polynucleotide is useful for diagnosing diseases caused by mutation in human NEDD-1, and for diagnosing or monitoring diseases caused by altered expression of human NEDD-1. Fragments of NEDD-1 are useful as hybridisation probes and primers, and to direct expression or synthesis of epitopic or immunogenic protein fragments. The proteins are useful as therapeutic supplement in patients with specific deficiency in human NEDD-1 production, and for treating subjects preferably with defects in NEDD-1. The present sequence is a PCR primer related to human NEDD-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HKNO1; ss; chromosome 18p; bipolar affective disorder; BAD; PCR; primer; severe bipolar affective (mood) disorder; BP-1; schizophrenia; Hong Kong new gene 1; antimanic; antidepressant; neuroleptic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        621 TAAGCTGGACAAACTGGGCG 640
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Gaps

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4; Indels

0; Mismatches

16; Conservative

Best Loca Matches

1684 TACATCTTCCCTGCTTACTC 1703

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recarcircereagnacic 20

ABN97923 standard; DNA; 20

RESULT 1349

ABN97923

ABN97923;

GAPDH amplification control forward primer. NEDD-1; cytostatic; human; ss; PCR; primer.

Homo sapiens

30-JUL-2002 (first entry)

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The invention relates to an isolated nucleic acid molecule comprising a nucleotide sequence that encodes a Hong Kong New Gene (HKNG) 1 gene product. The human gene for HKNG1 is located on chromosome 18p in an area associated with bipolar affective disorder, BhD. Also included are an expression vector comprising the nucleic acid, a host call expressing the nucleic acid, an anti-HKNG1 antibody, a method of identifying modulators of HKNG1, and identifying an individual (at risk of) having HKNG1 mediated disorder comprising detecting the presence or absence of a disorder, where the presence of the polymorphism indicates that the disorder at risk of having HKNG1 allele associated with the individual (is at risk of) having HKNG1—mediated disorder. A (small individual (is at risk of) having HKNG1—mediated disorder. A (small
                                                                                                                                                                                                                       New nucleic acid molecule Hong Kong New Gene 1 (HKNG1), useful for
screening for molecules which modulate HKNG1 expression for the treatment
of bipolar disorder and schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           molecule) compound which modulates (inhibits or potentiates) expression of a HKNG1 gene or gene product in a human individual is useful for the treatment of a HKNG1-mediated disorder with as bipolar affective disorder (BAD), severe bipolar affective (modd) disorder (BP-I) and schizophrenia. The present sequence is PCR primer which amplifies a HKNG1 exonic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Caspase 7; antisense modulation; antiinflammatory; cytostatic; antisense therapy; caspase 7 inhibitor; inflammatory condition; hyperproliferative disorder; cancer; bone metabolism; infection; cholesterol disorder; inflammation; tumour; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse caspase 7 phosphorothioate oligonucleotide SEQ ID NO:127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match
Local Similarity 80.0%; Pred. No. 9.3e+02;
es 16; Conservative 0; Mismatches 4; Indels
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/*tea b
/*mod base= OTHER
/note= "2'-methoxyethyl (2'-MOE) wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                               Disclosure, Page 74; 367pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             156 GTCAATGACACTCCGAGGTG 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 Grccardaaactrogadgrg 1
                                                                                                                                                   Novak T;
                                 02-AUG-2000; 2000US-00631275.
28-NOV-2000; 2000US-00722544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABN80949 standard; DNA; 20 BP
                                                                                       (MILL-) MILLENNIUM PHARM INC. (REGC ) UNIV CALIFORNIA.
 02-AUG-2001; 2001WO-US024417
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*tag= a
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                                                                                                                                               Chen H, Freimer NB,
                                                                                                                                                                                     WPI; 2002-195962/25
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modified_base
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Best Local Si
Matches 16
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ABN80949
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The present invention describes a compound (1) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding caspase 7, which specifically hybridises with and inhibits the expression of caspase 7.

(1) has antinifammatory and ofvostatic activities, and can be used in antisense therapy and as an inhibitor of caspase 7 expression. (1) is useful for inhibiting the expression of caspase 7 in human cells or tissues, and for treating a human having a disease or condition hyperproliferative disorder (cancer), or bone metabolism or cholesterol disorder. (1) is useful for diagnostics, therapeutics, prophylaxis and as research reagent and kits. (1) is useful prophylactically to prevent or delay infection, inflammation or tumour formation. The present sequence represent a mouse caspase 7 inhibiting chimeric phosphorothicate or represent a mouse caspase 7 inhibiting chimeric phosphorothicate
                                                                                                                                                                                                                                                                                                              Novel antisense compounds targeted to nucleic acids encoding caspase 7, for modulating gene expression and treating diseases associated with expression of caspase 7 in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Caspase 7; antisense modulation; antiinflammatory; cytostatic; antisense therapy; caspase 7 inhibitor; inflammatory condition; hyperproliferative disorder; cancer; bone metabolism; infection; cholesterol disorder; inflammation; tumour; phosphorothioate; see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Indels '0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.6; DB 1; Length 20; 30.0%; Pred. No. 9.3e+02; Ive 0; Mismatches 4; Indels
                /*tag= c
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        710 TCAGACTGGAACATGAAGAG 729
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                                                                                                                                                                              11-SEP-2000; 2000US-00659860.
                                                                                                                                                10-SEP-2001; 2001WO-US028232.
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Best Local Similarity 80.0%;
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                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                WPI; 2002-404806/43.
                                                                                                                                                                                                                                                Watt AT;
                                                                              WO200222640-A1
modified base
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modified_base
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                                                                                                              21-MAR-2002
                                                                                                                                                                                                                                              Zhang H,
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Homo sapiens.

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The present invention describes a compound (1) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding caspase 7, which specifically hybridises with and inhibits the expression of caspase 7.

(1) has antiinflammatory and cytostatic activities, and can be used in antisense therapy and as an inhibitor of caspase 7 expression. (1) is useful for inhibiting the expression of caspase 7 expression. (1) is useful for inhibiting the expression of caspase 7 in human cells or tissues, and for treating a human having a disease or condition associated with caspase 7 including inflammatory condition, hyperproliferative disorder (cancer), or bone metabolism or cholesterol disorder. (1) is useful for diagnostics, therapeutics, prophylaxis and as research reagent and kits. (1) is useful prophylactically to prevent or delay infection, inflammation or tumour formation. The present sequence oligonucleotide having 2. MOE wings and a decxy gap, which is used in an example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antisense compounds targeted to nucleic acids encoding caspase 7, for modulating gene expression and treating diseases associated with expression of caspase 7 in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human Von Willebrand factor-cleaving protease cloning PCR primer, 6395.
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80.0%; Pred. No. 9.3e+02;
ive 0; Mismatches 4; Indels
               1 . 5
/*tea b
/*mod base= OTHER
/note= "2'-methoxyethyl (2'-MOE) wing'
                                                                                                                                         'note = "2'-methoxyethyl (2'-MOE) wing"
'note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 8 A; 3 C; 7 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 88; 138pp; English.
                                                                                                                          base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 CTCTTTGCTTACTCCACGGT 1
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                                                                                                                                                                                                                                                                                 11-SEP-2000; 2000US-00659860,
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                                                                                   16. .20
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                                                                                                           /*tag=
                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-404806/43.
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Best Local Similarity
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                                                                                                                                                                           WO200222640-A1
               modified_base
                                                                                   modified_base
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Matches
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The invention relates to an isolated or substantially pure Von Willebrand factor-cleaving protease (vWF-cp) polypeptide. vWF-cp is useful for purifying vWF which involves providing vWF-cp as a ligand, contacting a solution comprising vWF with the polypeptide ligand under conditions where vWF is bound to the ligand and recovering from the ligand purified vWF. vWF-cp is useful for producing anti-vWF cp polypeptide antibodies which involves immunising an animal with vWF-cp and isolating the anti-vWF wF polypeptide antibodies from the animal. vWF-cp is useful for producing a preparation of prophylaxis and therapy of thrombosis and thrombombombolic disease such as thrombocic thrombocytopaenia or thrombombombolic disease such as thrombocic thrombocytopaenia or haemolytic-uraemic syndrome. vWF-cp can also be used for processing plasmatic or recombinantly produced vWF. The invention is useful for construction expression systems and generating transgenic animals which express the polypeptide in vivo. The present sequence is human vWF-cp
                                                                                                                                                                                                                                                                                                                                          Novel isolated or substantially purified Von Willebrand factor-cleaving protease, useful for producing preparation for therapy of thrombosis and thromboembolic disease such as thrombotic thrombocytic purpura.
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abnormal cell proliferation, neoplastic cell growth, growth-inhibitory.
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                                                                                                                                                                                                                  Lucumure B, Gerritsen HE, Furlan M, Turecek P, Schwarz
Scheiflinger F, Antoine G, Kerschbaumer R, Tagliavacca
Zimmermann K, Voelkel D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 6 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 34; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               253 CCTGGAGAGGCCCCCACACG 272
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                                                                                                        20-NOV-2001; 2001WO-EP013391.
                                                                                                                                           22-NOV-2000; 2000US-00721254
12-APR-2001; 2001US-00833328
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                                                                                                                                                                                                                                                                                                           WPI; 2002-479950/51.
                                                                                                                                                                                                 (BAXT ) BAXTER AG.
                                  WO200242441-A2.
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                                                                       30-MAY-2002.
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ABQ74705
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21-DEC-2001; 2001WO-US050574.

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The present invention describes a method for identifying a compound that induces senescence in a mammalian cell comprising culturing the cell in the presence and absence of the compound, assaying expression of at least con controllular gene (Glas) from 50 or agene (G2) from 64 genes, with corresponding accession numbers given in the specification, and identifying compounds that induce senescence when expression of (Glas) or expression of (G2) is lower, in the presence of the compound Also described: (1) a compound that induces senescence in a mammalian cell; (2) assessing efficacy of a treatment of a disease or condition relating to abnormal cell proliferation or neoplastic cell growth; or (4) identifying a compound that inhibits senescence—associated inducin of cellular gene expression or neoplastic cell growth; or (4) identifying a compound that inhibits is useful for treating to abnormal cell proliferation or senescence—associated induction of cellular gene expression or condition relating to abnormal cell proliferation or senescence—associated inducing the present of the invention has a growth—inhibitory effect without producing systemic side effects found with other growth-inhibitory compounds. ABQ74611 to ABQ74734 represent PCR primers which are used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; mouse; HYPLIP1; FCHL1; familial combined hyperlipidaemia; cancer;
                                                                                                                                                                                                                          Identifying a compound that induces senescence in a mammalian p53 deficient or tunor cell comprises assaying expression of cellular genes in the presence of the compound with expression of the genes in the absence of the compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP, 5 A; 3 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse HYPLIP1 locus PCR primer #302
                                                                                                                                                                                                                                                                                                                               Example 4; Page 52; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 48 ACCAGCAGTGTGACTGCTGA 67
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lipid disorder; PCR; primer; ss.
                                       21-DEC-2000; 2000US-0257907P.
17-DEC-2001; 2001US-00257907.
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                                                                                                                                                                                    WPI; 2002-619266/66.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified antisense oligonucleotide, antisense, HIV, cancer, infection, cytostatic, virucide, anti-HIV, hepatotropic, antiinflammatory, phosphorothioate backbone, integrin, cell-cell adhesion receptor, 8s.
                                                                                    New mouse HYPLIP1 and human FCHL1 (familial combined hyperlipidemia) genes and their sequence variations, useful for diagnosing, treating preventing lipid disorders and cancers.
                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                Lusis AJ;
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/note= "optionally phosphorothioate backbone"
11. .13
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                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                De Jong P,
                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
            Bodnar JS, Castellani LW, Chatterjee A, Ohmen J, Ross D, Tafuri S, Wu \in S_i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ICAM antisense oligonucleotide #1.
                                                                                                                                                Claim 11; Page 76; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                          16 GGATGGACAGGAATGCAGAG 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= c
/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 deargeagagacarcaga 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAL46755 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                        WPI; 2002-329882/36.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified
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The invention relates to a novel isolated nucleic acid molecule comprising a sequence that encodes a thioesterase or thioesterase domain,
                                                                                                                                                      The present invention relates to novel antisense oligonucleotides which are targetted to nucleic acids encoding human raf proteins and capable cinhibiting raf expression. The invention also relates to methods of inhibiting hyperproliferation of cells which involves contacting the hyperproliferating cells with a therapeutically effective amount of an oligonucleotide of the invention. The method is useful for treating cancer, angiogenesis or neovascularisation, especially ocular angiogenesis or neovascularisation. The present DNA sequence is an antisense oligonucleotide targetted to human c-raf kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated nucleic acid molecule from a bacterial daptomycin biosynthetic gene cluster encoding a thioesterase or thioesterase domain, useful for generating novel linear and cyclic peptides, and products in a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Daptomycin biosynthetic gene cluster; thioesterase; antibacterial; fungicide; virucide; antiparasitic; immunomodulator; antilipemic; cytostatic; gene therapy; antimitotic; immunomodulatory; siderophore; anti-cholesterolemic; agrochemical; linker; PCR; primer; ss.
                                                         Treating cancer, angiogenesis or neovascularization by administering antisense oligonucleotides targeted to human raf sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S. roseosporus daptomycin biosynthetic gene cluster PCR primer P92
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                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
vative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 91; 227pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1186 ATGGCCACAGGCCGTCCCCT 1205
                                                                                                                    Disclosure; Col 12; 41pp; English
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28-FEB-2001; 2001US-0272207P.
06-AUG-2001; 2001US-0310385P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABQ78911 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-OCT-2002 (first entry)
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Best Local Similarity 80.0
Matches 16; Conservative
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                       WPI; 2002-597918/64.
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(BALT/) BALTZ R H.
(SILV/) SILVA C J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200259322-A2.
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ABQ78911
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                                                                                                                                                                                                                                                                                         The present invention relates to oligonucleotides having at least one non-terminal pyrimidine nucleoside modified and additionally having the 5'-and/or 3'-terminal modified. These can be used in the treatment of viral infections, such as HIV, HSV-1, HSV-2, influenza virus, VSV, hepatitis B and papilloma viruses, cancer and diseases involving integrins and cell-adhesion receptors. The present sequence is an antisense oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; raf; hyperproliferation; neovascularisation; ocular angiogenesis;
therapy; cancer; cytostatic; anti-angiogenic; vascular; ophthalmological;
antisense; phosphorothioate backbone; c-raf kinase; ss.
                                                                                                 Winkler I;
                                                                                                                                                                           New nuclease-resistant oligonucleotides having modified non-terminal pyrimidine nucleoside(s), useful e.g. for treating cancer or viral diseases or as diagnostic reagents.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 13.6; DB 1; Length 20;
llarity 80.0%; Pred. No. 9.3e+02;
Conservative 0; Mismatches 4; Indels
                                                                                                 Helsberg M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human c-raf kinase antisense oligonucleotide ISIS #5149
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/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                   Kretschmar G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                           Disclosure; Page 12; 19pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GAGAGTGGTGGTGGTGGCGG 245
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95WO-US007111.
96US-00756806.
97US-00888982.
98WO-US013961.
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93DE-04338704
94EP-00117513
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*tag= a
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                                                                                                   Peymann A, Uhlmann E,
                                                                                                                                     WPI; 2002-353922/39.
                                                           (FARH ) HOECHST AG.
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16; Conserv
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modified_base
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Synthetic.
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12-NOV-1993;
07-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-1995;
26-NOV-1996;
07-JUL-1997;
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28-AUG-1998;
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RESULT 1357

AAD44724/

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Best Loca Matches

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Gaps

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derived from a bacterial daptomycin biosynthetic gene cluster. The proceins of the invention have antibacterial, fungicide, virucide, antiparasitic, immunomodulator, antilipemic, and cytostatic activity. The polynucleotides may have a use in gene therapy. The compositions and methods of the present invention are useful for generating novel linear and daptomycin non-ribosomal peptide synthetase (NRPS) to be used as new compounds or in producing new compounds, such as antibiotics, antifungals, antivirals, antiparasitics, antimitotics, antitumour agents, immunomodulatory agents, anti-cholesterolemic agents, siderophores, agrochemicals and cytostatics. The sequence represents a PCR primer used in the invention to amplify the S. roseosporus daptomycin biosynthetic gene cluster from a BAC library
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NOVX; cytostatic; cardiant; antiarteriosclerotic; antiasthmatic; cancer; hypotensive; cardiomyopathy; bronchial asthma; gene therapy; vaccine; human; PCR; primer; ss.
                                                                                                                                                                                                                                                                          Gaps
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0
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80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                       374 AGGCTTCAGCCACGTCCTCG 393
                                                                                                                                                                                                                                                                                                                        1 AGICCICAGCCAICICCICG 20
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2001US-0274134P.
2001US-027432P.
2001US-027432P.
2001US-0275235P.
2001US-0275235P.
2001US-0275235P.
2001US-0275239P.
2001US-0275239P.
2001US-0275239P.
2001US-0275239P.
2001US-0275331P.
2001US-0277331P.
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                                                                                                                                                                                                                                                                                                                                                                                                            ABX97255 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-MAY-2003 (first entry)
                                                                                                                                                                                                                                                           Best Local Similarity 80.0 Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200272757-A2.
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13-MAR-2001
13-MAR-2001
13-MAR-2001
14-MAR-2001
16-MAR-2001
19-MAR-2001
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-MAR-2001;
-MAR-2001;
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-MAR-2001;
-MAR-2001;
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PR 02-ARR-2001, 20018-028022P.

PR 02-ARR-2001, 20018-028032P.

PR 03-ARR-2001, 20018-028032P.

PR 15-ARR-2001, 20018-028032P.

PR 16-ARR-2001, 20018-028032P.
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0; Gaps

0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels

Query Match 0.8 Best Local Similarity 80.0 Matches 16; Conservative ô

The invention relates to a composition (I) comprising AGP-3 receptor (tumour necrosis factor ligand family member) related protein (II) attached to a vehicle protein. (I) is useful for modulating AGP-3-related activity in mesenteric lymph nodes (MLN) of a mammal. (II) is useful in assays to identify cells and tissues that express AGP-3R or proteins related to AGP-3R related protein and for identifying compounds (agonists or antagonists) that interact with AGP-3R proteins. (II) is also useful for identifying intracellular proteins that interact with the respective cytoplasmic domains by yeast two-hybrid screening process. (II) is anyolved in B cell growth, survival and activation particularly in lymph node, spleen, and Peyer's patches. AGP-3R agonists and antagonists involved using (II) are used for modulating B cell response and are used to treat diseases characterised by inflammatory processes or deregulated immune response such as rheumatoid arthritis, graft-versuschost disease. Crohn's disease, lupus, etc. (II) is also useful in the production of hybridoma cells which are derived from B cells, which involves treating the hybridoma cells with (II). (II) is useful in the transment of inflammatory conditions of joints or antagonists are useful for treating acute pancreatitis, and protophic lateral sclerosis (ALS), Alzheimer's disease, asthma, atherosclerosis, cachexia/anorexia, is schaemic injury including crebral ischaemia, multiple myeloma, multiple sclerosis, osteoporosis, Parkinson's disease, pain, reperfusion injury, dermatological, neuroprotective, nootropic; immunosmodulator; metabolic; antidiabetic; analgesic; nephrotropic; osteopathic; cytostatic; fever; antidiabetic; analgesic; nephrotropic; osteopathic; cytostatic; fever; antipartinsonian; antipartiatic; vasotropic; antibacterial; asthma; AGP-3 receptor; tumour necrosis factor ligand family; AGP-3 receptor; nesenteric lymph node; AGP-3R; inflammatory disease; immune disorder; rheumatorid arthritis; graft-versus-host disease; immune disorder; pancreatitis; amyotrophic lateral sclerosis; ALS; Alzheimer's disease; disease; glomerulonephritis; inflammatory bowel disease; ischaemia; ss; multiple sclerosis; Parkinson's disease; transgenic animal; PCR primer. Composition, useful for identifying modulator of receptor for treating asthma and glomerulonephritis, comprises AGP-3 (tumor necrosis factor ligand family member) receptor and encoding nucleic acids. Mouse; AGP-3; antiinflammatory; antiarthritic; immunosuppressive; Disclosure; Page 39; 124pp; English 306 CCCACTCAGCTCTGCACCAG 325 CCCATTCAGCACTGAAACAG 20 ВЪ. 12-FEB-2001; 2001WO-US004568 11-FEB-2000; 2000US-0181800P AAS18551 standard; DNA; 20 (first entry) Mouse AGP-3 PCR primer #5 WPI; 2002-049441/06. (AMGE-) AMGEN INC. Boyle WJ, Hau H; WO200185782-A2. Mus musculus. 12-MAR-2002 AAS18551; RESULT 1360 AAS18551/c δ

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Human; C/EBP beta; CCAAT/enhancer-binding protein beta; C/EBB2; LAP; TCF5; CRP2; NFIL6; IL6DBP; NF-M; AGP/EBP; Apc/EBP; transcription factor; tissue development; cellular function; proliferation; differentiation; hormone responsiveness; oxidative stress response;
IL-6 signalling mediator; interleukin-6; carbohydrate metabolism; famunity; Thi response; female fertility; gluconeogenesis; ovarian; cancer; tumour formation; type II; diabetes; infection; inflammation; expression inhibition; phosphorothioate; antisense oligonuclectide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE
cytosines are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE
cytosines are 5-methylcytosine"
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                                                                                                                                                                                                                                                                                                                                                                                     Human C/EBP beta phosphorothicate antisense oligonucleotide, SEQ ID:74
                                                                                                                                             Gaps
septic shock, etc. The nucleic acids are also useful for developing transgenic animals expressing (II), which are useful for producing the polypeptides and for the study of in vivo biological activity. The present sequence represents mouse AGP-3 PCR primer #5
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                                                                                                         0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
iive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "Phosphorothioate linkages"
                                                                            Sequence 20 BP; 6 A; 7 C; 7 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                           ABL94308 standard; DNA; 20 BP
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                                                                                                             Query Match 0.89
Best Local Similarity 80.09
Matches 16; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                           ABL94308;
                                                                                                                                                                                                                                                              RESULT 1361
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Example 8; Page 65; 81pp; Japanese.

of e.g. tumor.

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cothe human or mouse CCAAT/enhancer-binding protein alpha (C/EBP alpha)

cothe human or mouse CCAAT/enhancer-binding protein alpha (C/EBP alpha)

designed to target different regions of the human and/or mouse C/EBP

alpha RNA, and were analysed for their effect on C/EBP alpha mRNA levels

by quantitative real-time PCR. The C/EBP family of proteins are a family

of transcription factors which regulate the expression of a wide range of

genes that control normal tissue development, cellular function, cellular

cofferation and functional differentiation. C/EBP beta (also known as

c/EBB2, LAP, TGES, CRE2, NRIE6, IL6DBP, NB-M, AGP/EBP and Apc/EBP)

primarily regulates hormone responsiveness and oxidative stress responses

and is a mediator of IL-6 (interleukin-6) signalling. C/EBP beta is

thought to be involved in carbohydrate metabolism, immunity, the Th1

response, female fertility and gluconeogenic pathways. C/EBP beta is

cropped to the liver lung, spleen, kidney, brain, and testis, with the

lighest expression in pancreas is upregulated in response to chronically

and its expression in pancreas is upregulated in response to chronically

clevel in malignant ovarian tissue compared with normal ovarian tissue,

and its expression in pancreas is upregulated in response to chronically

clevel the invention are useful for diagnosis, prevention and treatment of

conditions associated with C/EBP beta expression, such as cancer

(particularly ovarian cancer), tumour formation, diabetes (particularly

type II diabetes), infection, or inflammation
Sequences ABL94252-ABL94476 represent antisense oligonucleotides targeted to the human or mouse CCAAT/enhancer-binding protein alpha (C/EBP alpha)
          8KGGGGGGGGGGGGGGGGGGGGGGGGG
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Sequence 20 BP; 2 A; 8 C; 7 G; 3 T; 0 U; 0 Other;

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4; Indels 0;
ch 0.8%; Score 13.6; DB 1; Length 20; 1 Similarity 80.0%; Pred. No. 9.3e+02; 16; Conservative 0; Mismatches 4; Indels
                                                                                     65 TGAAACCCAGGGGAGGGCCC 84
      Query Match
Best Local S
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20 TGAGACTCCGGGGAGCGCC 1 g

ABK49114 standard; DNA; 20 BP. 02-JUL-2002 (first entry) ABK49114; RESULT 1362 ABK49114

Human KDR/FLK-1 mutagenic PCR primer for Y801F mutant.

Human; KDR; kinase insert domain-containing receptor; FLK-1; ss; fetal liver kinase-1; cytostatic; antidiabetic; antirheumatic; antiarthritic; signal transduction; phosphorylation; cell proliferation; anglogenesis; tumour; diabetic omentopathy; chronic rheumatoid arthritis; PCR; primer; mutant.

Homo sapiens.

Synthetic.

WO200229090-A1.

02-OCT-2001; 2001WO-JP008684. 11-APR-2002.

03-OCT-2000; 2000JP-00303694. (KYOW) KYOWA HAKKO KOGYO KK. (SHIB/) SHIBUYA M. Shibuya M, Takahashi T, Furuya A, Shitara K;

WPI; 2002-352237/38.

Screening substances inhibiting the binding of signal-transducing molecule to KDR/Flk-1 phosphorylated at tyrosine at 1175-position, as cell proliferation inhibitors and angiogenesis inhibitors for treatment

The invention relates to inhibiting the signal transduction of KDR/FIk-1 (Kinase insert domain-containing receptor/fecal liver kinase-1) is by using a substance inhibiting the binding of a signal-transducing molecule to KDR/FIk-1 phosphorylated at tyrosine at the 1175-position. Also included are methods of detecting/inhibiting/screening for cell conjuded are methods of detecting/inhibiting/screening for cell complete at tyrosine at the 1175-position. Also inhibitors, compounds obtained by the screening methods, drugs containing the inhibitors, compounds obtained by the screening methods, drugs containing to the inhibitors, a monoclonal antibody or its fragment, a recombinant vector containing the monoclonal antibody or its fragment, a recombinant vector containing the conclonal antibody or its fragment, a recombinant vector containing the phosphorylated at tyrosine at the 1175-position, a DNA encoding the containing the binding of a signal-transducing molecule to KDR/FIk-1 conhibiting the binding of a signal-transducing molecule to KDR/FIk-1 conhibitors and anglogenesis inhibitors for treatment of e.g. tumour, diabetic omentopathy and chronic rheumatoid arthritis. A method for cettering anglogenesis is also provided. The present sequence is a PCR to Phe

Sequence 20 BP; 3 A; 6 C; 5 G; 6 T; 0 U; 0 Other;

0; Gaps Match
 0.8%; Score 13.6; DB 1; Length 20;
Local Similarity 80.0%; Pred. No. 9.3e+02;
ies 16; Conservative 0; Mismatches 4; Indels Query Match

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RESULT 1363

ABI97222 standard; DNA; 20 ABI 97222;

BP.

(first entry) 16-FEB-2002

Capture oligonucleptide Zip ID#4309 oligo #9.

Human, K-ras, PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.

Synthetic.

WO200179548-A2.

25-OCT-2001.

04-APR-2001; 2001WO-US010958.

14-APR-2000; 2000US-0197271P.

(CORR) CORNELL RES FOUND INC.

Gerry NP, Favis R, Kliman Barany F, Zirvi M,

MPI; 2002-034366/04.

Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.

Example 5; Fig 29; 300pp; English

%XGGGGGGGGGGGGGGGGGGGGGGGGG

The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents (e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergilus fundsautus, viruses e.g. T-cell lymphocytorrophis cirus, Epstein-Barx virus and policy of T. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, psi gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and federity and compresses scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe processes course of the target nucleotide sequences. ABIS2074 to the target nucleotide sequences in the exemplification of the present invention

Sequence 20 BP; 7 A; 4 C; 7 G; 2 T; 0 U; 0 Other;

0; Gaps Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels

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AAS20906 standard; DNA; 20 BP. AAS20906/c CXSXLLLXSXLXBXBXBXBXBXXBXBXBXBXBXCXC

(first entry) 09-APR-2002 AAS20906;

Human peptide transporter hPHT1 cDNA RT-PCR primer #3.

Human; peptide histidine transporter 1; hPHT1; peptide transport; peptide-based drug transport; cell membrane; gastrointestinal tract; hPHT1-related disease; reverse transcriptase; RT-PCR; primer; ss.

WO200192468-A2. Homo sapiens

06-DEC-2001.

31-MAY-2001; 2001WO-US017650.

31-MAY-2000; 2000US-0208061P.

(RUTF) UNIV RUIGERS STATE NEW JERSEY.

Knipp GT, Herrera-Ruiz D;

WPI; 2002~130529/17.

Novel isolated human peptide histidine transporter which facilitates peptide transport across cell membranes in gastrointestinal tract, useful as target for evaluating peptide and peptide-based drug transport.

Example 2; Page 55; 95pp; English.

The present invention relates to nucleic acid sequences encoding human

peptide histidine transporter 1 (hPHT1) protein, the hPHT1 proteins and methods for using them. The nucleic acid sequences of the invention are is useful for screening a test compound for human PHT1 modulating cativity. The hPHT1 proteins are useful as a target for evaluating cativity. The hPHT1 proteins are useful as a target for evaluating peptide and peptide-based drug transport. The functional characterisation of the hPHT1 proteins are useful as a target for a particular substrate to the molar expression level of hPHT1 provides crucial information regarding the ability of this transporter to facilitate the uptake and transport across cell membranes in the PHT1 proteins facilitate peptide transport across cell membranes in the pHT1 protein facilitate and other organs in which they are expressed. The identification of full length hPHT1 close facilitates the development of optimal peptide-based drugs for treating patients with hPHT1-related diseases. AAS20818-AAS20811 represent reverse transcriptase (RT)-PCR primers used in the methods of the present invention

Sequence 20 BP; 0 A; 5 C; 12 G; 3 T; 0 U; 0 Other;

Gaps ·. 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ative 0; Mismatches 4; Indels 16; Conservative Best Local Similarity Query Match Matches

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RESULT 1365 ABK67749

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ABK67749 standard; DNA; 20

ABK67749;

02-JUL-2002 (first entry)

Mouse transglutaminase associated PCR primer #9.

Transglutaminase; TGM; transamidation; autoimmune disease; Addison's disease; Al heamolytic anaemia; Al thrombocytopenic purpara; Al thyroid disease; Al heamolytic anaemia; Al thrombocytopenic purpara; Al thyroid disease; atrophic gastritis; pernicious anaemia; Chron's disease; colitis ulcerosa; Goodpasture syndrome; IgA nephropathy; Egg diomerulonephritis; mysathenia gravis; partial lipodystrophy; polymyositis; primary biliary cirthosis; primary sclerosing cholangitis; progressive systemic sclerosis; recurrent pericarditis; primary claromic pericarditis; systemic atrophy; diabetes; Wegener granulomatism; sarcoidosis; SLE; splenic atrophy; diabetes; Wegener granulomatosis; ulcerative colitis; vasculitis; vitiligo; PCR; primer; ss.

Mus sp

WO200222830-A2.

21-MAR-2002.

14-SEP-2001; 2001WO-GB004120.

15-SEP-2000; 2000GB-00022768. 16-MAY-2001; 2001GB-00011995.

(UYCA-) UNIV COLLEGE CARDIFF.

Grenard PM; Aeschlimann DP,

WPI; 2002-329954/36.

and TG-Y Nucleic acids which encode novel transglutaminase enzymes TG-Z which can be used in diagnostic methods of autoimmune diseases.

Disclosure, Page 27; 67pp; English.

The invention relates to nucleic acids which encode noval polypeptides having transglutaminase activity. The compositions of polypeptide are useful for transamidation reactions on peptides and polypeptides.

Detection of the polypeptides with transglutaminase activity are useful in a subject or in calls derived from a subject to a diagnostic method in a subject or in calls derived from a subject by a diagnostic method in a subject or in calls derived from a subject proteins may be used to diagnose autoimmune diseases which include the proteins may be used to diagnose autoimmune diseases which include they cold diseases, a trophic gastrictis, permicious anaemia, Chron's disease, colitis ulcerosa, Goodpasture syndrome, IgA nephropathy or IgG glomerulonephritis, mysathenia gravis, partial lipodystrophy, progressive systemic sclerosis, recurrent pericarditis, relapsing proparties, primary biliary dirthosis, primary sclerosing cholangitis, polymyositis, prematoid arthritis, rheumatism, sarcoidosis, Sjogren's syndrome, SiE, splenic atrophy, type I (insulin-dependent) diabetes mellitus, Wegener granulomatosis, ulcerative colitis, vasculitis (both syndrome cutaneous) and vitiligo. This sequence represents a primer used in the study of transglutaminase genes in which DNA, amino acid sequence. determined 88888888888888888888888

Seguence 20 BP; 5 A; 3 C; 9 G; 3 T; 0 U; 0 Other;

Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels 599 TTGGGAAACTGGAGACCTAC 618 ò

rrecedacrecadedadecade 20

AB081403;

ABQ81403 standard; DNA; 20 BP

Arabidopsis AINTEGUMENTA-like gene PCR primer.

(first entry)

12-DEC-2002

Lipid metabolism regulator; LTR; plant; transgenic plant; transcription factor; seed oil; oilseed; cardiant; wril; AINTEGUMENTA; PCR; primer; ss.

Arabidopsis thaliana.

WO200272775-A2.

19-SEP-2002

08-MAR-2002; 2002WO-US007441.

D8-MAR-2001; 2001US-0274170P

(BADI) BASF PLANT SCI GMBH.

Benning C, Cernac A;

WPI; 2002-713509/77.

New isolated lipid metabolism regulator nucleic acid, useful for producing transgent plants having modified level of seed storage compound, e.g. lipids for generating seed oils which have the ability of reducing risk of heart disease.

Example 2; Page 34; 72pp; English.

The present sequence is that of a primer for an AINTEGUMENTA-like protein gene of Arabidopsis thallana. Overlapping PCR primers (see ARG1398-407) were used in amplification and sequencing reactions to identify sequence changes in 2 wril mutants compared to wild-type sequences in order to identify the true wril gene. In subsequent experiments, wril mutants were complemented with cosmids containing wild-type genomic DNA, and PCR was used to produce a full-length wril CDNA (see ARG81395) encoding a lipid RESULT 13
RESULT

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metabolism regulator (LMR) protein (see ABB79954). LMR is suggested to act as a transcription factor regulating lipid and seed storage compound metabolism during seed development. The invention relates to the use of LMR nucleic acids in the production of transgenic plants having a modified level of a seed storage compound. The level of a lipid, fatty acid, starch or seed storage protein can be modified, yielding a seed obtaint is medically and nutritionally useful in reducing the risk of heart
                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                         Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02;
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0.8%; Score 13.6; D
Best Local Similarity 80.0%; Pred. No. 9.3e
Matches 16; Conservative 0; Mismatches
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Human Mac2-BP promoter PCR primer SEQ ID NO: 68. ABT08433 standard; DNA; 20 (first entry) 27-NOV-2002 ABT08433; RESULT 1367 ABT08433

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Gaps ;

Human, cyclin-dependent kinase, CDK, cyclin-dependent kinase inhibitor; inhibitor; cancer; age-related disease; promoter; atherosclerosis; cytostatic, antiarteriosclerotic; nootropic; neuroprotective; nephrotropic; antiarthritic; arthritis; renal disease; Alzheimer's disease; amyloidosis; PCR; primer; ss.

Homo sapiens.

WO200266681-A2.

29-AUG-2002.

01-FEB-2002; 2002WO-US002784.

01-FEB-2001; 2001US-0265840P. 21-MAY-2001; 2001US-00861925.

(UNII) UNIV ILLINOIS FOUND.

Chang Poole J, Roninson IB,

m ;

WPI; 2002-674960/72.

New recombinant expression construct, useful for identifying compounds that inhibit the induction of genes induced by cyclin-dependent kinase inhibitors for preventing or treating cancer, renal failure or Alzheimer's disease.

Example 11; Page 133; 137pp; English.

The present invention relates to a recombinant expression construct encoding a reporter gene operably linked to a promoter from a mammalian gene induced by a cyclin-dependent kinase (CDK) inhibitor. The construct is useful for identifying compounds that inhibit the induction of genes induced by CDK inhibitors. The compounds are useful for preventing or tracting a disease caused by CDK inhibitor induced gene expression, e.g. cancer other than colon cancer, renal failure, Alzheimer's disease, amyloidosis, age-related diseases, atherosoclerobis or arthritis. The present sequence is a PCR primer used to amplify a human promoter suitable for use in the construct of the invention

Sequence 20 BP; 5 A; 3 C; 7.G; 5 T; 0 U; 0 Other;

schultz621-3.rng

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The invention relates to a method of measuring beta 1,4-N-acetylgalactosaminyltransferase (GD2/GM3 synthase) mRNA. The method involves obtaining an mRNA sample, performing real-time quantitative reverse transcriptase-polymerase chain reaction (RT-PCR) on the sample using appropriate primers of GD2 synthase, and determining the amount of GD2 mRNA. The methods and kits are useful for detecting and/or diagnosing various forms of cancer such as neuroblastoma, melanoma, B cell lymphoma, osteosarcoma, soft tissue sarcoma, medullablastoma, high-grade astrocytoma, retinoblastoma, Wilm's tumour, Ewing's sarcoma, bladder carcinoma, lung cancer, breast cancer, pancreatic cancer, oseophageal cancer, pastrointestinal cancer, sarcoma, head and neck tumours or melanoma. The present sequence is BACE marker gene specific RT-PCR primer, used to illustrate the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                        Measuring GD2 synthase mRNA, useful for detecting or diagnosing cancer, e.g. neuroblastoma, small cell lung cancer, melanoma, by performing realtime quantitative RT-PCR on the sample using appropriate primers of GD2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, bifunctional apoptosis regulator, antisense, phosphorothioate,
cytostatic, antiinflammatory; inhibitor, infection, inflammation, tumour,
    cancer; neuroblastoma; melanoma; lymphoma; carcinoma; sarcoma; tumour;
primer; BAGB; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human bifunctional apoptosis regulator antisense oligo ISIS NO 143737
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80.0%; Pred. No. 9.3e+02;
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                                                                                                                                                                                                                                                                                                          Cheung NV;
                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-129279/12.
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modified_base
                                                            Unidentified.
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                                                                                                                                         21-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABX78206,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel recombinant blood coagulation factor VIII, its production process and its medicinal composite for treating type-A haemophilia. The invention further comprises a medicinal composition containing the blood coagulation factor which promotes blood coagulation to the blood plasma of type-A haemophilia patients. This polynucleotide sequence represents an oligo relating to the recombinant blood coagulation factor VIII protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New recombinant blood coagulation factor VIII and its production process and medicinal composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beta 1, 4-N-acetylgalactosaminyltransferase; GD2 synthase; GM2; RT-PCR; reverse transcriptase PCR; medullablastoma; astrocytoma; retinoblastoma;
                                           Gaps
                                                                                                                                                                                                                                                                                                                                               Recombinant blood coagulation factor VIII protein related oligo #11.
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ch 0.8%; Score 13.6; DB 1; Length 20; Similarity 80.0%; Pred. No. 9.3e+02; 16; Conservative 0; Mismatches 4; Indels
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80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                        blood coagulation factor VIII; type-A haemophilia; ss.
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                                                                                48 ACCAGCAGTGTGACTGCTGA 67
                                                                                                                     1 Accargacicide 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-DEC-2000; 2000CN-00137779.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-DEC-2000; 2000CN-00137779
                                                                                                                                                                                                                           ADE64605 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD53075 standard; DNA; 20
                                                                                                                                                                                                                                                                                                     29-JAN-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-741852/81.
Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Qi Z, Wang Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
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                                                                                                                                                                              RESULT 1368
ADE64605/c
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Matches
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AAD53075
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23-APR-2002; 2002WO-US013135.

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This invention describes a novel compound, 17-50 nucleobases in length which specifically hybridises with a nucleic acid encoding human bitch specifically hybridises with a nucleic acid encoding human bitch solutions appoptosis regulator (BAR) and inhibits the expression of human BAR. The products of the invention have cytostatic and antiinflammatory activity and can be used to inhibiting the expression of during antisense therapy, useful for inhibiting the expression of human contains an animal, particularly a human suspected of having expression of BAR in an animal, particularly a human suspected of having or being prone to a disease or condition associated with expression of human baR. In addition the antisense oligonaclecities are useful for diagnostics, therapeutics and as research reagent, e.g. prophylactically to prevent or delay infection, inflammation or tumor formation. The coligonuclecities described in the invention human BAR antisense oligonuclecities and a deoxy gap. This sequence represents a human BAR antisense oligonuclecities described in the disclosure of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, antisense, lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
  5
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-14 are 2'-deoxy- nucleotides, all C nucleotides are methyl cytosines"
                                                                                                                                                                                                                                                                                                                      New antisense compounds targeted to nucleic acids encoding human bifunctional apoptosis regulator, for modulating expression of the regulator and treating diseases associated with expression of the regulator in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ve 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Col 45-46; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   195 CAATGGTGCCCTGAGCAGA 214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 caardccarccraagaga 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human oligonucleotide sequence
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                                                                                                                                                                      27-APR-2001; 2001US-00844525.
                                                                                                                                 27-APR-2001; 2001US-00844525
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nes 16; Conservative
                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                   WPI; 2003-196749/19
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                                                          US6468796-B1
                                                                                               22-OCT-2002
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Matches
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initialion coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of installation codon, coding region, 5' or regions within 2-10 mucleotides of chast all airway dysfunctions and a second active agent comprising an antiinflammatory steroid and ubjoinform. A composition of the invention has antiinflammatory antiallargic, antiasthmatic, hypotensive, as a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of adenosine of or reducing sensitivity to adenosine, reducing levels of ubjquinone or lung surfactant in a subject, to treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                           Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; broncholilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
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                                                                                                                                              Pabalan J, Aguilar D;
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0
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Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 4 A; 9 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 5692; 872pp; English.
                                                                                                                                              Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1034 ACTITGGCCTGGCCCGAGCC 1053
                                                                                                                                              Li Y, Sandrasagra A, K
Tang L, Shahabuddin S;
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                                                             24-APR-2001; 2001US-0286137P.
                                                                                                     (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ92603 standard; DNA; 20
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                                                                                                                                                                                                       WPI; 2003-229219/22.
                                                                                                                                              Ľį Y,
                                                                                                                                              Nyce JW, ]
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WO200285308-A2 Homo sapiens

31-OCT-2002.

31-OCT-2002

schultz621-3.rng

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Pharmaceutical composition for treating ailments associated with impaired
                                                                                              respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid
                                                                                                                          Disclosure; SEQ ID NO 7845; 872pp; English.
       23-APR-2002; 2002WO-US013135.
                    24-APR-2001; 2001US-0286137P.
                                   (EPIG-) EPIGENESIS PHARM INC
                                                                        WPI; 2003-229219/22.
                                                                                                            biquinone.
                                                 Nyce JW, I
Miller S,
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of Junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubsquinone. A composition of the invention has mutinflammatory, antiallergic, antiasthematic, hypetensive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also cor enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubsquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition.

Specification, but was obtained in electronic format directly from WIPO
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0.8%; Score 13.6; DB 1; Length 20; 0.0%; Pred. No. 9.3e+02; ve 0; Mismatches 4; Indels 80.08; Similarity Query Match Best Local S

16 GGATGGACAGGAATGCAGAG 35 GGATGGCCGGGACTGCACAG 1 ઠે

16, Conservative

Matches

ABZ88825 standard; DNA; 20 BP ABZ88825; ABZ88825

(first entry) 17-0CT-2003

Human oligonucleotide sequence.

Human, antisense; lung dysfunction, nasal airway dysfunction, antiinflammatory; antiallergic; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytoetatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, K Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-229219/22.

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Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar D; Tang L, Shahabuddin S;

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid ubiquinone

Disclosure; SEQ ID NO 4067; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intiation cooden, coding region, 5' or 3' end genemic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal alrawy dysfunction and a second active agent comprising an antiminamatory steroid and ubjudinton. A composition of the invention has antiminamatory, antiallergic, antiasthmatic, hypotensive, communication subjects of an antiminamatory antiallergic, antiasthmatic, hypotensive, as use in antistense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, as 180 or canding sensitivity to adenosine, reducing or depleting levels of antiminamatory steroid in a subject, for reducing or depleting levels of a denosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubjquinome or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allamation, lung allamation, shear is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 6 A; 4 C; 1 G; 9 T; 0 U; 0 Other;

Gaps ð., 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indel8 Query Match 0.8 Best Local Similarity 80.0 Matches 16; Conservative

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Gaps

; 0

RESULT 1374 ABZ87133/c

ABZ87133 standard; DNA; 20 BP

ABZ87133;

(first entry) 17-0CT-2003 Human oligonucleotide sequence.

antiinflammatory steroid, ubiquinone, antiinflammatory, antiallergic, antiasthmatic, hypotensive; immunosuppressive, cytostatic, gene therapy, antisense gene therapy; respiratory; lung, adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. antisense; lung dysfunction; nasal airway dysfunction;

Homo sapiens

WO200285308-A2 BX3X8X244444

31-OCT-2002

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first active agent comprising an interpolation and intration codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encodings a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an intilifiammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antisflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchocanstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this parent is not represented in the printed of specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel pharmaceutical composition, which has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 15; SEQ ID NO 2375; 872pp; English.
                                                                                                                                                                                                                                                                                          Li Y, Sandrasagra A, K
Tang L, Shahabuddin S;
23-APR-2002; 2002WO-US013135.
                                                                                               24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                               (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-229219/22.
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                                                                                                                                                                                                                                                                                               Nyce JW,
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0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels Sequence 20 BP; 2 A; 7 C; 6 G; 5 T; 0 U; 0 Other; 1285 GCCATCCTGTCCAACGAGGA 1304 Query Match
Best Local Similarity 80.0
Matches 16; Conservative à

20 GGCATCCGGACCAGCGATGA 1

ABZ92417 standard; DNA; 20 BP (first entry) 17-0CT-2003 ABZ92417; RESULT 1375 BXXXXXXXXXXXXXXXXXX

Human oligonucleotide sequence,

Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory steroid, ubiquinone, antiinflammatory, antiallergic, antiasthmatic; hypotensive, immunosuppressive, cytostatic, gene therapy, antisense gene therapy, respiratory; lung; adenosine sensitivity, adenosine receptor; bronchodilation, bronchoconstriction; lung allergy; lung inflammation, respiratory disease, ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Katz E, Pabalan J, Li Y, Sandrasagra A, K, , Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pabalan J, Aguilar

Katz E,

Pharmaceutical composition for treating allments associated with impaired or respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid ubiquinone.

Disclosure, SEQ ID NO 7659; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligomucleotide antisense to the intitation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an artification and associated with lung and/or nasal airway dysfunction and biquinone. A composition of the invention has mutinflammatory antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a use in antisers gene therapy. The composition may have a use in antisers are piratory, lung or malignant disease or condition, also preventing a respiratory, lung or malignant disease or condition, as a continulammatory steroid in a subject, for reducing or depleting levels of or enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of adenosable receptor, producing bronchodilation, increasing levels of adenosable receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition.

Supecification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 5 A; 9 C; 4 G; 2 T; 0 U; 0 Other;

Gaps ö Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02; 0; Mismatches 4; Indels 80.08; Query Match 0.8° Best Local Similarity 80.0 Matches 16; Conservative

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Gaps ö

ABZ88076 standard; DNA; 20 BP ABZ88076; RESULT 1376 ABZ88076

Human oligonucleotide sequence. (first entry) 17-0CT-2003

Human, antisense; lung dysfunction, nasal airway dysfunction, antiinflammatory steroid, ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytosteatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchodilation; bronchodilation; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens,

WO200285308-A2.

31-OCT-2002.

schultz621-3.rng

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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                     Li Y, Sandrasagra A, Katz B, Pabalan J, Aguilar D;
Tang L, Shahabuddin S;
       23-APR-2002; 2002WO-US013135.
                       24-APR-2001; 2001US-0286137P.
                                      (EPIG-) EPIGENESIS PHARM INC
                                                                              WPI; 2003-229219/22.
                                                     Nyce JW, I
Miller S,
                                                                                                                     ubiquinone.
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first active agent comprising an oligonucleotide antisense to the intration codon, coding reggion, 5' or 3' end genomic flanking regions, initiation codon, coding reggion, 5' or 3' end genomic flanking regions, intran-exon junctions or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of denosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject, for tracting bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed of specification, but was obtained in electronic format directly from WIPO Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other; at ftp.wipo.int/pub/published_pct_sequences Disclosure; SEQ ID NO 3318; 872pp; English.

0; Gaps 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ve. 0; Mismatches 4; Indels Best Local Similarity 80.0%; Matches 16; Conservative Query Match ઠે

457 GAGGACATCAACAAGCGCCT 476 GAGGAGCTCAACAAGCTGCT 20

ABZ88262 standard; DNA; 20 BP ABZ88262; ABZ88262/

(first entry) 17-0CT-2003 Human oligonucleotide sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction, antinflammatory steroid, ubiquinone, antinflammatory; antiallergic, antiasthmatic; hypotensive, immunosuppressive; cytostatic; gene therapy, antisense gene therapy; respiratory; lung, adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(BPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, K Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 3504; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosition inflammation, lung allergies, or a respiratory disease or condition, or the printed producine but was obtained in electronic format directly from WIPO specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 7 A; 6 C; 3 G; 4 T; 0 U; 0 Other;

ö Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02; 0; Mismatches 4; Indels 0.8%; Ouery Match 0.8 Best Local Similarity 80.0 Matches 16; Conservative

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RESULT 1378 ABZ84865

ABZ84865 standard; DNA; 20 BP. ABZ84865; BXBXSXEXEXEXEXEXBXBXB

17-OCT-2003 (first entry)

Human oligonucleotide sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory steroid; ubiquinone, antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy, antisense gene therapy, respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens,

WO200285308-A2

31-OCT-2002.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the intraraction codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubjquinone, A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytotatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also containing sensitivity to adenosine, reducing levels of an antiinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodiation, increasing levels of adenosine receptor, producing bronchodiation, increasing levels of ubjquinone or lung surfactant in a subject sissue, or treating bronchoconstriction, lung surfactant in a subject sissue, or treating bronchoconstriction, lung surfactant in a subject sissue, or treating bronchoconstriction, lung surfactant in a subject sissue, or treating bronchoconstriction, lung surfactant down as obstanced in electronic format directly from WIPO

Specification, but was obtained in electronic format directly from WIPO

The treation of the producing potential processor of the printed services.
                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating allments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                   Li Y, Sandrasagra A, Katz E, Pabalan J,
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 15; SEQ ID NO 107; 872pp; English.
                            23-APR-2002; 2002WO-US013135.
                                                                            24-APR-2001; 2001US-0286137P.
                                                                                                                               (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                               WPI; 2003-229219/22.
                                                                                                                                                                                     Nyce Jw, I
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                     ubiquinone.
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Gaps .. 0 Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels Sequence 20 BP; 3 A; 11 C; 2 G; 4 T; 0 U; 0 Other;

299 CACGGGGCCCACTCAGCTCT 318 cacrerceccacecagerer 20

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ABZ85601/c ID ABZ85601 standard; DNA; 20 BP ABZ85601; RESULT 1379

17-OCT-2003 (first entry)

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic; antiinflammatory; antiallergic; antianflamtinatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

VO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Katz E, Pabalan J, Nyce JW, I Miller S,

Aguilar D;

Li Y, Sandrasagra A, K Tang L, Shahabuddin S;

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Claim 15; SEQ ID NO 843; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flamking regions, 5' or 3' and 3' intron-exon junctions, 5' or so regions within 2-10 nuclectides of junctions of gense encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensis of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensis or composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or mallignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, bronchodilation, increasing bronchoconstriction, lung antifarmation, lung allergies, or a respiratory disease or condition.

Specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published_pct_sequences

Seguence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;

Gaps ö Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels

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0

RESULT 1380

ABZ86435 standard; DNA; 20 BP ABZ86435;

17-OCT-2003 (first entry)

Human oligonucleotide sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory steroid; ubiquinone, antiinflammatory, antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy, antisense gene therapy, respiratory, lung, adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2

31-OCT-2002.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5 or 3 end genomic flanking regions, 5 and 3 intron-exon junctions, or regions within 2-10 nuclectides of Junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubsquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, and cytostatic activity. The composition may have a use in antiinflammatory, antiallergic, antiasthmatic, hypotensive, composition and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also preventing a respiratory, lung or malignant disease or condition, also of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition.

Specification, but was obtained in electronic format directly from MPPO
                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                              respiration, has oligo(s) antisense to specific gene(s) or its
corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 5 A; 2 C; 6 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                     Claim 15; SEQ ID NO 1677; 872pp; English
                                                                                                                                   Li Y, Sandrasagra A,
Tang L, Shahabuddin
                   23-APR-2002; 2002WO-US013135.
                                                          24-APR-2001; 2001US-0286137P.
                                                                                                (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ92850 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                    WPI; 2003-229219/22.
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                                                                                                                                                                                                                                                                                                     ubiquinone.
                                                                                                                                     Nyce JW,
Miller S,
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Matches
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ö Gaps .. 0 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels 1481 TCCACAAACTTCCTGACACT 1500 16; Conservative Local Similarity

20 TCCAGAAACGTCTTAACACT 1

ВЪ

Human oligonucleotide sequence.

antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; de. Human; antisense; lung dysfunction; nasal airway dysfunction;

Ното варіепв.

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar D;

Pabalan J,

Katz E, S;

Aguilar D; Pabalan J, Katz E, ŝ Sandrasagra A, L, Shahabuddin Li Y, San , Tang L, Miller S, Nyce JW,

WPI; 2003-229219/22.

treating ailments associated with impaired or respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid Pharmaceutical composition for ubiquinone

Disclosure; SEQ ID NO 8092; 872pp; English.

intiation relates to a movel phratmaceutical composition, which has a initiation codon, coding region, 5' or 3' end genomic flanking regions, 6' or 3' end genomic flanking regions within 2-10 mucleotides of increase ancoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory staroid and ubiquinone. A composition of the invention of the antiinflammatory, antiallergic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an infilammatory staroid in a subject, for reducing or depleting levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition.

So precification, but was obtained in electronic format directly from WIPO or security in the printed or security. invention relates to a novel pharmaceutical composition, which has a at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 4 A; 3 C; 4 G; 9 T; 0 U; 0 Other;

Gaps ö Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02; 0; Mismatches 4; Indels 0.8%; Query Match Best Local Similarity 80.0 Matches 16; Conservative

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ABZ75967 standard; DNA; 20 ABZ75967; RESULT 1382 ABZ75967/

BP.

(first entry) 29-MAY-2003

ICAM-1 gene targeting 2'-deoxyoligonucleotide ISIS 1939.

ICAM-1; desulphurization; antioxidant; intercellular adhesion molecule-1;

Homo sapiens Synthetic

WO2003005822-A1

23-JAN-2003

11-JUL-2002; 2002WO-US022038

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Hormone-sensitive lipase, antisense oligonucleotide, inhibitor; obesity; phosphorochioace, antidabetic; anorectic; cytostatic; antisense therapy, abnormal metabolic condition; hyperlipidaemia; type 2 diabetes; cancer; hyperproliferative disorder; human; ss.
                                                                                                                                                                                                                 The invention relates to preventing desulphurization of an oligonuclectide or its bioequivalent comprising one or more phosphorothicate linkages in a bi-phasic or multi-phasic formulation. The method involves including in the formulation an anticoxidant which partitions into the aqueous phase of the formulation. The method is useful for increasing the stability of oligonucleotide comprising phosphorothicate linkages. The present sequence represents a 2-decoxyoligonucleotide having a phosphorothicate backbone and is targeted to the 3' UTR (untranslated region) of ICAM-1 (intercellular adhesion
                                                                                                                         Preventing desulfurization of oligonucleotide comprising phosphorothioate linkages in bi-phasic/multi-phasic formulation, by adding to formulation an antioxidant that partitions into aqueous phase of the formulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human HSL chimeric phosphorothioate oligonucleotide SEQ ID NO:106.
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                      Disclosure; Page 12; 51pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           226 GAGAGTGGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 GAGAGGGGAAGTGGTGGGGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JUL-2002; 2002WO-US022672.
 11-JUL-2001; 2001US-00902953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ82717 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                              (ISIS-) ISIS PHARM INC
                                                                                            WPI; 2003-229426/22
                                                              Mehta R;
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modified_base
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                                                              Krotz AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ82717;
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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/note= "2'-O-methoxyethyl (2'-MOE) wing"

mod_base= OTHER

υ *tag=

mod_base= OTHER note= "2'-O-methoxyethyl (2'-MOE) 16. .20

1. .5 /*tag= b

New antisense oligonucleotides targeted against nucleic acids encoding hormone-sensitive lipase, useful for treating abnormal metabolic condition, e.g. hyperprolipidemia and obesity, or a hyperproliferative disorder, e.g. cancer. Gaps Inhibition, phosphorothioate, delayed release oral formulation; phosphorothioate, delayed release oral formulation; rheumatoid gastrointestinal absorption; ulcerative colitis; rheumatoid arthritis; Crohn's disease; inflammatory bowel disease; abnormal cellular proliferation; ss. . Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02; 0; Mismatches 4; Indels ICAM-1 inhibitory antisense oligonucleotide SEQ ID NO:2. /note= "phosphorothioate linkages" Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other; Wyatt JR; Example 15; Page 89; 167pp; English. Location/Qualifiers 1003 ATCAACGAGAGAGAGCT 1022 /*tag= a /mod base= OTHER Freier SM, ATCACCGAGATGGAAGTGCT 1 170/c ACC49170 standard; DNA; 20 BP 26-JUL-2001; 2001US-00915814. Query Match
Best Local Similarity 80.0%;
Matches 16; Conservative 22-AUG-2002; 2002WO-US026924 (first entry) (ISIS-) ISIS PHARM INC Watt AT, WPI; 2003-239411/23. WO2003017940-A2 Key modified_base 19-JUN-2003 06-MAR-2003 Synthetic. Butler MM, 20 ACC49170; RESULT 1384 ACC49170/c à 유

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The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding a hormone-sensitive lipase (HSL) or a splice variant of HSL. The compound specifically hybridises with an least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding HSL. (I) have anorectic, antidabetic and cytostatic activities, and can be used in antisense therapy. (I) is useful for treating an animal, particularly human, suspected of having an abnormal metabolic condition such is obesity, hypertipidaemia, type 2 diabetes, a hyperprisiferative disorder such as cancer (e.g. pituitary, colorectal, breast, testicular, pulmonary or epithelial cancer). (I) is also useful in modulating blood glucose levels particularly plasma or serum glucose levels, in a diabetic animal. The present sequence represents a human hormone-sensitive lipase chimeric phosphorothioate antisense oligonucleotide, which is used in an example from the present invention
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Gaps

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3 TT:0T:40 7004
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(ISIS-) ISIS PHARM INC

Geary RS, Hardee GE, Tillman LG, Weinbach SP,

WPI; 2003-354422/33.

Pulsed release oral formulation providing enhanced gastrointestinal absorption, comprises first particles containing drug and penetration enhancer and second particles containing delayed release penetration

Disclosure; Page 28; 59pp; English.

The present invention describes a delayed release oral formulation (A), giving enhanced gastrointestinal (GI) absorption of a drug (I). (A) comparises a first set of particles containing (I) and a penetration comparises a first set of particles containing (II) in a delayed release coating or matrix (III). (A) is used for enhancing the absorption of [I] in mammals, especially humans. Typical disorders to be treated include ulcerative colitis, rheumatoid arthritis, Crohn's disease, inflammatory bowel disease and abnormal cellular proliferation. When the particles release (I) and (II) at a first location in the GI tract clease pulse) and is often present in insufficient amount to promote absorption of the entire dose of (I). This problem is solved by providing further (II) in delayed release form in the particles, so that absorption of (I) is completed in a second buse. The present sequence represents an exemplary oligonucleotide from the present invention which inhibits ICAM-

Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;

0; Gaps Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 9.3e+02; les 16; Conservative 0; Mismatches 4; Indels Query Match Best Local S: Matches 16

226 GAGAGTGGTGGTGGCGG 245

20 GAGAGGGAAGTGGTGGGGG 1

163/c ACC62163 standard; DNA; 20 BP

ACC62163;

20-JUN-2003 (first entry)

Human alipoprotein B antisense oligonucleotide SEQ ID NO: 52.

alipoprotein B; ApoB; antilipaemic; antiarteriosclerotic; antidiabetic; anorectic; cardiovascular; gene therapy; lipid metabolism; cholesterol metabolism; atherosclerosis; hyperlipidaemia; diabetes; type 2 diabetes; obesity; atherosclerosis; cardiovascular disease; glucose; antisense oligonucleotide; ss.

Synthetic

WO2003011887-A2.

13-FEB-2003.

30-JUL-2002; 2002WO-US024247.

01-AUG-2001; 2001US-00920033. 30-APR-2002; 2002US-00135985. 15-MAY-2002; 2002US-00147196.

(ISIS-) ISIS PHARM INC

Graham MJ;

WPI; 2003-268105/26.

New antisense oligonuclectides for modulating apolipoprotein B, especially for preventing or treating atherosclerosis, hyperlipidemia or diabetes, or for modulating glucose, cholesterol, lipoprotein or triglyceride levels.

Example 15; Page 96; 160pp; English.

The invention relates to a novel compound that is 8-50 nucleotides in length that is targeted to a nucleic acid molecule encoding and proportion and inhibits appropriate and ancleic acid molecule encoding ApoB; or which the expression of a nucleic acid molecule encoding ApoB; or which specifically hybridises with at least an 8-nucleotide portion of an active site on a nucleic acid molecule encoding ApoB; or which active site on a nucleic acid molecule encoding ApoB; or which active site on a nucleic acid molecule encoding ApoB; or which active site on a nucleic acid molecule encoding ApoB; or which active site on a nucleic acid molecule encoding ApoB; or which an activity and satisface of specifically proportion associated with ApoB; e.g. a condition involving abnormal lipid metabolism, a condition involving an abnormal metabolism, atherosclerosis or a condition involving an abnormal metabolism, atherosclerosis or a condition and annual metabolism, atherosclerosis or activity an abnormal metabolism, atherosclerosis or cardiovascular disease). The new compound or the antisense cardiovascular disease). The new compound or the antisense oligonucleotide is also useful for modulating glucose levels (specifically vabL, MDL or LDL) or serum triglyceride levels canimal or for modulating serum cholesterol levels, lipoprotein levels (specifically vabL, MDL or LDL) or serum triglyceride levels or particularly in a human. The antisense compound is also useful for the new compound is also useful for the new or disease or condition associated with ApoBs, or the onset of a disease or condition to the the mimmal or human. The present sequence is used in the exemplification of the invention

Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

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0; Gaps Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels

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ABX13023 standard; DNA; 20 BP.

ABX13023;

10-MAY-2003 (first entry)

Oxidative stress detection PCR primer #64.

Oxidative stress detection; PCR; primer; ss; risk factor

Homo sapiens.

WO2003016527-A2.

27-FEB-2003.

13-AUG-2002; 2002WO-EP009079.

14-AUG-2001; 2001BE-00000545.

(PROB-) PROBIOX SA.

Crooke RM,

BCDUITZ621-3.rnd

enhancer #X#X#X#X#####X#X#####

22-AUG-2001; 2001US-00944493

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ACC62163,

ä Pincemail J, Piette J, Marechal

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Determining oxidative stress markers in a group of individuals by comparing the amount of each of the oxidative stress markers obtained from each of the group of individuals with that of the group of healthy individuals.
                                                                                                                                                                                                                                                                    Disclosure; Page 36; 67pp; English.
WPI; 2003-268334/26
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markers in a group of individuals. The method comprises determining the risk factor for oxidative stress in the group, measuring the amount of at least 10 different oxidative stress markers in a sample obtained from each of the group of individuals, and comparing the amount of each of the oxidative stress markers with the amount of each of the oxidative stress markers with the amount of each of the oxidative stress the oxidative stress markers are increased or decreased in the group of individuals carrying a risk factor for oxidative stress relative to healthy individuals. This sequence represents a PCR primer used to detect The invention relates to a method for determining oxidative stress oxidative stress

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Sequence 20 BP; 5 A; 3 C; 7 G; 5 T; 0 U; 0 Other;
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Gaps .; 0 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; 4; Indels 0; Mismatches Conservative Similarity Query Match Best Local Simil Matches 16; C

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621 TAAGCTGGACAAACTGGGCG 640
                                   resectreschaderes 20
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ABX33984 standard; DNA; 20 (first entry) 10-FEB-2003 ABX33984; RESULT 1387 ABX33984/c g

BP

Human interleukin 12 p40 subunit antisense oligonucleotide ISIS #139157. Human, ss, antisense; interleukin 12 p40 subunit, antibacterial, antiinflammatory; cytostatic; infection; inflammation; tumour. Homo sapiens

/note= "All cytosines are 5-methylcytidines and the nucleotides are linked via a phosphorothicate backbone" Location/Qualifiers 1. .20 /*tag= a /mod_base= OTHER /*tag= b /mod_base= OTHER Key modified_base modified_base modified base

US6448081-B1

10-SEP-2002

07-MAY-2001;

07-MAY-2001; 2001US-00851062.

(ISIS-) ISIS PHARM INC

Freier SM; Baker BF,

WPI; 2003-074100/07.

New antisense chimeric oligonucleotide, useful for modulating the expression of human Interleukin 12 p40 subunit, in treating or preventing disease states in humans and animals, and as research reagents and diagnostics.

Example 15; Col 45; 42pp; English

length targeted to a start codon region, coding region, a stop codon region or a 3'-untranslated region of a nucleic acid molecule encoding human Interleukin 12 p40 subunit. The compound specifically hybridises with one of the regions and inhibits the expression of human Interleukin 12 p40 subunit. The new compound is useful for inhibiting the expression of human Interleukin 12 p40 subunit in cells or tissues and comprises contacting the cells or tissues in vitro with the compound, so that antisense compound may also be used as research reagents and diagnostics, and as treatment or prevention of disease states, e.g. to prevent or delay infection, inflammation or tumour formation, in animals and humans The present sequence is an antisense oligonucleotide of the invention The invention relates to an antisense compound 20-50 nucleobases in

Sequence 20 BP; 4 A; 4 C; 9 G; 3 T; 0 U; 0 Other;

Gaps ö Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02; 0; Mismatches 4; Indels 0.8%; Ouery Match Best Local Similarity 80.0 Matches 16; Conservative

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ABZ83986 standard; DNA; 20 1388 ABZ83986 RESULT

BP. (first entry) 14-MAY-2003 ABZ83986;

Toxicologically relevant rat PCR primer #1145.

Toxicologically relevant gene; toxicological response; PCR primer; ss.

Rattus sp. Synthetic.

WO2003016500-A2

27-FEB-2003.

16-AUG-2002; 2002WO-US026514

(PHAS-) PHASE-1 MOLECULAR TOXICOLOGY INC. 16-AUG-2001; 2001US-0313080P.

Schmeiser Kier LD, Pickett GG, Adkins K, Neft RE, Dunn RT, Alen P;

WPI; 2003-268322/26.

Determining a toxicological response to an agent, useful for screening of drugs, comprises comparing the expression profile of one or more human toxic response genes to a reference gene expression profile indicative of

Claim 1, Page 326, 455pp, English.

The present invention describes a method (M1) for determining

schultz621-3.rng

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toxicological response to an agent, which comprises comparing the expression profile of one or more human toxic response genes to a reference gene expression profile indicative of toxicity, and so determining the presence of a toxic response to the agent. Also described: (1) an array comprising one or more polynucleotides selected from the genes corresponding to the partial sequences given in ABS2842 to ABS84764, or their fragments of at least 20 nucleotides, or homologues; and (2) determining if a gene putatively identified to be a toxic response gene playes a role on toxic response pathways by determining the expression profile of the gene after exposure of cells or a human subject to a known toxic pharmaceutical or industrial agent, comprising: (a) expossing cells to an agent or isolating cells from a human subject to a known toxic pharmaceutical or industrial agent, and (c) comparing the test or a putatively identified toxic response gene after exposure to a known toxic pharmaceutical or industrial agent; and (c) comparing the test profile to the expression profile of a gene with a similar function or comparing the test profile to the expression profile of a gene with a similar function or comparing the test profile to the expression profile of a gene with a similar function or comparing the test profile to the expression profile of a gene with a similar function or predicting and determining toxic compounds. The methods are useful for predicting and determining toxicological responses on a cellular, organ or system level. The arrays compiraling the human genes are useful for toxic lostice and a cellular, organ or system level. The arrays compiration the human genes are useful for toxicological screening of drugs, pharmaceutical compounds and chemicals
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Sequence 20 BP; 5 A; 2 C; 10 G; 3 T; 0 U; 0 Other;

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Gaps
/ Match 0.8%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 9.3e+02; Loss 16; Conservative 0; Mismatches 4; Indels
                                                                                           357 TGATGGGGAGAGTGACCAGG 376
       Query Match
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ADA26797 standard; DNA; 20 BP ADA26797; RESULT 1389 ADA26797/

20-NOV-2003 (first entry)

Human PRL-3 forward PCR primer #81.

Metastasis; neoplastic growth; detection; prediction; neoplastic growth marker; drug screening; cancer; tumour; gastrointestinal; prostate; breast, colorectal; diagnostic imaging; drug targeting; chromosome 8Q24.3; human; protein tyrosine phosphatase type IVA member 3; PRL-3; cytostatic; reverse transcription-PCR; RT-PCR; primer; ss.

Homo sapiens

WO2003031930-A2. 17-APR-2003. 02-OCT-2002; 2002WO-US031247.

09-OCT-2001; 2001US-0327332P.

(UYJO) UNIV JOHNS HOPKINS.

Vogelstein B, Kinzler KW, Saha S, Bardelli A; 4PI; 2003-393457/37. Identifying regions of neoplastic growth in a human body, useful for detecting or predicting metastasis, comprises administering to the human body an antibody or peptide that specifically binds to a protein marker of neoplastic growth.

Example 2; Page 22; 42pp; English

The invention relates to methods for identifying regions of neoplastic growth in a human patient, especially for detecting or predicting or metasates. The methods involve determining whether a neoplastic growth marker protein is overexpressed, either by the use of an antibody specific for the protein, or by the use of FCR or hybridisation to detect nucleic acids encoding the marker proteins. A set of neoplastic growth markers are disclosed (SAGE (serial analysis of gene expression) tags for these are given in ADA26759-ADA26796), with protein tyrosine phosphatase type IVA member 3 (also known as FRL-3) being a preferred neoplastic growth marker. The neoplastic growth markers are specifically expressed at a higher level in metastatic cancers. Compared with advanced and early growth markers are specifically expressed at a higher level in metastatic cancers. The invention also oversypression of the neoplastic growth markers is taken as an indication oversypression of the neoplastic growth markers is taken as an indication that the tissue has a propensity to metastaise. The invention also cancer, and for identifying candidate drugs for treating advanced or metastatic cancers. The methods of the invention are useful for identifying regions of neoplastic growth, for detecting or predicting metastasis, or identifying candidate drugs for treating advanced or metastatic cancers. The invention is particularly applicable to metastatic cancers. The invention is particularly applicable to metastatic cancers. The invention is particularly applicable to metastatic imaging and for targeting orthocial cancers. Antibodies which bind to the neoplastic growth marker proteins are additionally useful for diagnostic imaging and for targeting orthocial cancers. Antibodies them has present sequence represents a reverse them are presented at chromosome 8q24.3) in an example of the invention of the present sequence or supplementary applicable. invention. 8KGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG

Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;

0; Gaps / Match
0.8%; Score 13.6; DB 1; Length 20;
Local Similarity 80.0%; Pred. No. 9.3e+02;
les 16; Conservative 0; Mismatches 4; Indels Query Match

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ACD42082 standard; DNA; 20 RESULT 1390

05-SEP-2003 (first entry) ACD42082;

Antisense oligonucleotide targeting human c-raf, ISIS5149.

Human; 88; antisense; c-raf; a-raf; b-raf; protein kinase; cancer; signal transduction; cell proliferation; lung carcinoma; cytostatic; antisense gene therapy; chemotherapeutic agent; angiogenesis; hyperproliferative condition; neovascularisation; ocular angiogenesis.

Homo sapiens.

US2003032607-A1.

13-FEB-2003

31-MAY-1994; 94US-00250856. 31-MAY-1995; 95WO-18007111. 26-MOV-1996; 9TUS-00756806. 07-UUL-1997; 9TUS-00888982. 06-UUL-1998; 9BWO-US013961. 18-BAUG-1998; 9US-00143214. 25-JAN-2002; 2002US-00057550

(MONI/) MONIA B P.

Monia BP;

Erlich HA, Bugawan TL, Noble JA, Valdez AM;

Mirel

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The invention relates to an oligonucleotide 8-50 nucleotides in length which is targeted to mRNA encoding human c-raf, a-raf or b-raf (raf is a protein kinase playing a regulatory role in signal transduction. regulating cell proliferation and which is capable of inhibiting the expression. Also included is a composition comprising the oligonucleotide and a pharmaceutically acceptable carrier. The antisense oligonucleotide is useful for inhibiting the expression of human raf in human cells or issues, by contacting the human cells or tissues with the oligo. The oligo. is also is useful for treating or preventing a disease or condition with a chemotherapeutic agent to a human or cells of the human, where the expression of raf by administering it in combination with a chemotherapeutic agent to a human or cells of the human, where the expression of raf is abnormal expression, and the condition is a neovascularisation (preferably coular angiogenesis or neovascularisation). The oligo, is also useful for inhibiting hardered example for detecting and determining the role of raf expression in various cell functions and abhalological processes and conditions and physical processes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present sequence is an antisense oligonucleotide targeting
                                                                                        Novel antisense oligonucleotide which is targeted to mRNA encoding human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnosing conditions associated with raf expression and for research purposes. The present sequence is an antisense oliconucleotide target
                                                                                                                                             raf and which is capable of inhibiting raf expression, useful for
treating or preventing hyperproliferative conditions such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.6; DB 1; Length 20; ilarity 80.0%; Pred. No. 9.3e+02; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                    Disclosure; Page 7; 42pp; English.
WPI; 2003-503332/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human raf mRNA
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Matches
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ABQ80152 standard; DNA; 20 BP (first entry) 13-JUN-2003 ABQ80152; RESULT 1391

Right primer DBM0071B amplifies IL4R amplicon of 177 bp.

Human; interleukin 4 receptor; IL4R; type 1; diabetes; allele; insulin dependent diabetes mellitus; IDDM; wasthenia gravis; PCR; single nucleotide polymorphism; SNP; autoimmune disease; amplify; T helper type 1 mediated disease, rheumatoid arthritis; primer; multiple sclerosis; inflammatory bowel disease; systemic sclerosis; eystemic lupus erythematosus; psoriasis; scleroderma; grave's disease; guillain-Barre syndrome; Hashimoto's thyroiditis; se.

Homo sapiens.

WO2003010335-A2

06-FEB-2003

17-JUL-2002; 2002WO-EP007956.

20-JUL-2001; 2001US-0306912P.

(HOFF) ROCHE DIAGNOSTICS GMBH. (HOFF) HOFFMANN LA ROCHE & CO AG F.

The sequences given in ABQ80141-52 represent primers which were used to identify wild type and variant loci in the human interleukin 4 receptor CC (ILAR). These primer sequences were used in the method of the invention of determining an individual's risk for type 1 diabetes. The method comprises detecting the presence of an insulin dependent diabetes comprises detecting the presence of an insulin dependent diabetes comprises detecting the presence of the allele in a nucleic acid sample of the individual, where the presence of the allele indicates the individual's risk for type 1 diabetes. The method identifies one or comprises in the fort type 1 diabetes are useful for determining an individual's risk for any autoimmune disease or condition or any T helper type 1 mediated disease, systemic lupus cor any T helper type 1 mediated disease, systemic lupus cor any T helper type 1 mediated disease, systemic lupus cor any T helper type 1 mediated disease, systemic lupus corpusing myasthenia gravis, gclerosis, scleroderma, Grave's disease, systemic type in thy colditis comprises detecting Determining an individual's risk for type 1 diabetes, comprises detecting the presence of an insulin dependent diabetes mellitus-associated interleukin 4 receptor allele in a nucleic acid sample of the individual. Example 4; Page 35; 79pp; English. WPI; 2003-248086/24.

4; Indels 0.8%; Score 13.6; DB 1; 80.0%; Pred. No. 9.3e+02; iive 0; Mismatches 4; GGAGATTCAGCTACAAAAGG 1540 Query Match
Best Local Similarity 80.0
Matches 16; Conservative 1521 ઠે

1 gcadacrcadcaacaadage 20

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Gaps

Sequence 20 BP; 8 A; 5 C; 6 G; 1 T; 0 U; 0 Other;

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Gaps

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Length 20;

ACC49159 standard; DNA; 20 BP ACC49159,

ACC49159;

(first entry) 19-JUN-2003 ICAM-1 inhibitory antisense oligonucleotide SEQ ID NO:2.

Inhibition, antisense oligonucleotide; phosphorothioate; bioadhesive; enhanced mucosal drug absorption; antiulcer; antinflammatory; cancer; antirheumatic; antiarthritic; cytostatic; ulcerative colitis; tumour; rheumatoid arthris; Oroh's disease; inflammatory bowel disease; cellular proliferation; ss.

Synthetic

/note= "phosphorothioate linkages" Location/Qualifiers base= OTHER ø . .20 *tag= /mod Key modified_base

WO2003018134-A2

06-MAR-2003.

22-AUG-2001, 2001US-00935316.

22-AUG-2002; 2002WO-US026925.

(ISIS-) ISIS PHARM INC

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The present invention describes an oral pharmaceutical formulation (I) for delivering a bioreactive macromolecule to a mucosal surface. (I) comprising a pipulation of carrier particles comprising drug and a bioaddesive compound; and a second population of carrier particles comprising a penetration enhancer. Also described is a method for enhancing the mucosal absorption of the bioactive macromolecule in a mammal (preferably a human) by mucosally administering (I). (I) has artilloer, antiliflammatory, antilheumatic, antiarthritic and cytostatic activities. (I) can be used for delivering a bioreactive macromolecule to a mucosal surface. It is used for the oral delivery of a drug to an animal encompassing a human as well as other mammals, reptiles, fish, amphibians and birds. It is used to deliver drugs including peptiles, proteins, monoclonal antibodies their fragments, nucleic acids (DNA and RNA), oligonucleotides, antisense oligonucleotides, and small molecules. It can be used to examine the function of various proteins and genes in an animal, including those that are sessential to animal development. It can be used for the treatment of animals that are known or suspected to suffer from any disease treatable with the inventive composition, e.g. ulcerative collits, rheumatodid arthritis, crobin's disease, inflammatory obeyel disease, or undue cellular proliferation (cancers and tumours). The present sequence represents an exemplary oligonucleotide from the present invention, which can be used to inhibit ICAM-1
                                                                                       Oral pharmaceutical formulation for delivering bioreactive macromolecule to mucosal surface, contains drug, bioadhesive compound, and penetration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR; primer; Vpr; ss; immune response; immunocompromise; HIV; cancer; gene therapy.
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80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                Weinbach SP, Tillman LG, Geary RS, Hardee GE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                        Disclosure; Page 28; 62pp; English.
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                                                        WPI; 2003-342432/32.
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(TANN/) TANNER J.
(ROUX/) ROUX P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
Matches 16; Conserv
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enhancer.
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Roux

Alfieri C, Tanner J,

Gaps

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The invention relates to a DNA or RNA construct capable of expressing interleukin (IL)-2 in a warm-blooded animal or biological preparation, comprising a Vpr activated promoter, a transcribable DNA segment coding for a signal peptide functional in mammary cells and operably linked between the promoter and the DNA segment to facilitate secretion of IL-2. The construct is useful for increasing the immune response of a warm-blooded animal or biological preparation, by introducing the construct in stem cells, antigen preparation, by introducing the construct in stem cells, antigen corresponse of a warm-blooded animal or biological preparation to obtain a calls, of the warm-blooded animal or biological preparation. The warm-blooded animal is an immal or biological preparation. The warm-blooded animal is an immunocompromised patients. The method is useful for stimulating immune response in immunocompromised patients. The present sequence represents and diven construct associated primer. Note: The present sequence is made to it is the sequence listing but no further reference is made to it
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                                              Novel DNA or RNA construct for increasing immune response of warm-blooded animal, has Vpr activated promoter, DNA segment encoding interleukin 2 and secretory DNA encoding signal peptide functional in mammary cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotide, cytostatic; immunosuppressive; antiinflammatory; gene therapy; hyperproliferative disorder; cancer; autoimmune; inflammatory disorder; inhibitor-kappa B kinase-gamma; ss;
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/note= "Phosphorothioate linkages, all cytosines are
methylcytosine"
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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. 20
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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
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                                                                                                                         Disclosure; Page 15; 28pp; English.
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modified_base
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The specification describes a method of analysing DNA or RNA targets. The method comprises contacting the targets with oligonucleotide probes attached to an enzyme cofactor marker which is recognized less by the enzyme when it is on a free oligonucleotide than when it is on a hybridized oligonucleotide. The method is useful for analysing DNA or RNA targets. The invention is used to determine the amount of a target nucleic acid in a biological sample and the level of complementarity between the probe and the target nucleic acid. The present sequence represents an oligonucleotide probe attached to flavin. an enzyme cofactor marker for flavin reductase. The probe, together with its complement ABZ77540, was used to study the activity of flavin reductase
                                                                                                                                                                                    Analyzing DNA or RNA targets, useful for determining nucleic acid in a
biological sample, comprises using probes marked by a cofactor for an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                      (COMS ) COMMISSARIAT ENERGIE ATOMIQUE.
(UYFO-) UNIV FOURIER JOSEPH.
                                                                                                     Dueymes C;
                                                                                                                                                                                                                                                                         Example 1; Page 15; 36pp; French.
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16-JUL-2001; 2001FR-00009460.
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Best Local Similarity 80.0%;
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                                                                                                       Decout JL,
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                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an antisense compound that is targeted to a nucleic acid encoding inhibitor-kappa B Kinase-gamma, specifically hybridising to the nucleic acid encoding inhibitor-kappa B Kinase-gamma and inhibiting its expression. Compounds of the invention are antisense cligonucleotides comprising at least one modified internucleoside invention are antisense cligonucleotides comprothioate linkage, at least one modified sugar modety, which is a 2-0-methoxyethyl sugar modety, or at least one modified nucleobase, which is a 5-methylcytosine. Preferably, the modified nucleobase, which is a 5-methylcytosine. Preferably, the invention is useful for preparing a composition for treating a hyperproliferative disorder e.g., cancer, or an autoimmune or hyperproliferative disorder e.g., cancer, or an autoimmune or inflibitor-kappa B Kinase-gamma in cells or tissues, and treating an animal having a disease or condition associated with thibitor-kappa B kinase-gamma. Sequences given in ADA44713-ADA44790 represent antisense oligonucleotides for the inhibition of human mRNA levels.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                  New compound having sequence targeted to nucleic acid encoding inhibitor-kappa B kinase-gamma, useful for preparing composition for treating e.g., cancer, or inflammatory or autoimmune disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Probe; enzyme cofactor marker; enzyme; flavin; flavin reductase; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleotide seguence of a probe for flavin reductase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 8 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/note= "-p-C6flavin attached"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                        Example 15; Page 77; 106pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               917 TGTTCCTGTTCCAGCTGCTC 936
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                                                                                                     06-OCT-2001; 2001US-00972607.
                                                          03-OCT-2002; 2002WO-US031809.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABZ77539 Standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 80.0 les 16; Conservative
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                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                        Monia BP, Wyatt JR;
                                                                                                                                                                                                                              WPI; 2003-457242/43.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FR2827304-A1
                      17-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  substrate; ligand; signal; ligand binding; immobilisation;
gene engineering; genetic engineering; structure; biological activity;
ligand-receptor binding; PCR primer; amplification; ss.
                                                                     Gaps
                                                                     .
0
Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
                                                                 4; Indels
                                                                 0; Mismatches
                                                                                                                                   226 GAGAGTGGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           p38 gene PCR primer SEQ ID NO:22.
                                                                                                                                                                                              20 cacaccaactccicccc
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Example 1; Page 33; 52pp; Japanese.

The present invention describes a substrate having a number of ligands which have been immobilised onto a predetermined region of its surface, in which the region on the substrate has such a shape as to allow the concentration of signals caused by binding of the ligands to receptors in the region toward the receiver. Also described is a substrate for the immobilisation of such ligands. The substrates are applicable in general information of such ligands. The substrates are applicable in generalization of such ligands. The substrates and biological activity e.g. effect of endocrine disrupters on various genes and also in investigating the effect of hormones, drugs and other chemicals on the environment. Such substrates are highly sensitive in detecting the ligand-receptor binding, with affinity and reproducibility. The ligandimmobilised substrates can be produced in high density e.g. in microarray form to provide finely tuned results. ADA00221 to ADA00282 represent PCR prometry. Sequence 20 BP; 5 A; 5 C; 3 G; 7 T; 0 U; 0 Other; invention

Gарв ö Length 20; 4; Indels 0.8%; Score 13.6; DB 1; 80.0%; Pred. No. 9.3e+02; iive 0; Mismatches 4; Local Similarity 80.0 les 16; Conservative Query Match Matches

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ઠ 셤 RESULT 1397

ABZ23813 standard; DNA; 20 BP.

ABZ23813;

18-MAR-2003

EGFR mRNA inhibiting antisense oligonucleotide AS3. (first entry)

Epidermal growth factor receptor; EGFR; cytostatic; cancer; EGF antisense; ss

Synthetic

Homo sapiens.

WO200290514-A2.

14-NOV-2002

07-MAY-2002; 2002WO-US014557

07-MAY-2001; 2001US-0289055P.

(HYBR-) HYBRIDON INC.

Agrawal S, Kandimalla ER;

WPI; 2003-120540/11.

New synthetic oligonucleotide complementary to nucleic acids encoding epidermal growth factor receptor (EGFR), useful for inhibiting the EGFR gene or mRNA expression, and reducing cancer cell proliferation.

Claim 10, Page 12; 36pp; English.

complementary to a region of nucleic acid encoding epidermal growth factor receptor (EGFR) with location 245-1117, 2407-3201, 3786-4102 or 4574-45633. The methods and compositions of the invention are useful for annancing inhibition of EGFR gene or mRNA expression, and reducing cancer cell proliferation, in particular cancer cells of the colon, ovarian or breast. Sequences ABZ23811-832 represent specific examples of such The invention relates to synthetic antisense oligonucleotides

ö Gaps antisense oligonucleotides that inhibit the EGFR mRNA expression ö 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; 4; Indels Sequence 20 BP; 2 A; 4 C; 7 G; 7 T; 0 U; 0 Other; 0; Mismatches 1553 GGTCTTCGTCGATGCCTGAC 1572 16; Conservative Local Similarity Query Match Best Loca Matches ខ្លង់ខ្ល ਨੋ

1 Gercirgércaarerciege 20

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RESULT 1398

ABX78149 standard; DNA; 20 BP

ABX78149;

16-APR-2003 (first entry)

Murine p38-alpha MAPK antisense oligonucleotide ISIS NO 100812.

p38 mitogen-activated protein kinase, p38 MAPK; phosphorothioate, antisense; antiarthritic; antiinflammatory; kinase inhibitor; mouse; inflammatory disease; rheumatoid arthritis; gene therapy; ss.

Mus musculus

Location/Qualifiers Key modified_base

1. .20 /*tag= a /mod_base= OTHER

/note= "nucleotides 1-5 & 16-20 are 2'-methoxyethoxy (MOE) nucleotides, nucleotides 1-4 * 16 10

(MOE) nucleotides, nucleotides 1-4 & 16-19 are linked via phosphodiester linkages, nucleotides 6-15 are 2'-deoxy-nucleotides, nucleotides 5-16 are linked via phosphorohioate linkages, all C nucleotides are 5-methyl cytosines"

US6448079-B1

15-AUG-2000; 2000US-00640101.

99US-00286904 06-APR-1999;

(ISIS-) ISIS PHARM INC

Mckay Gaarde WA, Nero P, Monia BP,

ä

WPI; 2003-089122/08.

New antisense compound, useful for preparing a composition for diagnosing, treating or preventing inflammatory diseases, e.g. rheumatoid arthritis.

Example 5; Col 27-28; 44pp; English

This invention describes a novel antisense compound, which is 8-30 mucleobases in length targeted to a nucleic acid molecule encoding p38 mitogen-activated protein kinase (MAPK). He products of the invention have antiarthitic and antihilammatory activity, can act as act as kinase inhibitors. The antisense compound is useful for preparing a composition for diagnosing, treating or preventing inflammatory diseases, e.g. rheumatoid arthritis or for use in antisense gene therapy. This sequence represents an antisense oligonucleotide used in a method to inhibit p38 MAPK

Sequence 20 BP; 6 A; 7 C; 5 G; 2 T; 0 U; 0 Other;

Query Match

DB 1; Length 20; 0.8%; Score 13.6; schultz621-3.rng

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ABT43268;
                                                                                                                                                            Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences ABZ74952-ABZ74991 represent antieense oligonucleotides targeted to the human p70 S6 kinase gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human p70 S6 kinase RNA, and were analysed for their effect on mRNA levels by quantitative real-time PCR. p70 S6 kinase (also known as SK6, p70/p85 s6 kinase, p70/p85 ribosomal S6 kinase and pp70s6k) is a serinethreonine kinase responsible for the phosphorylation of the ribosomal S6
                                                                                                                                                                                                                                                                                                                                                              /note= "Phosphorothioate linkages. When bases 1-5 and 16-20 are not 2'-methoxyethyl (2'-MOB) nucleotides, all cytosines in the oligonucleotide are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antisense compound which is targeted to nucleic acid encoding p70 S6 kinase, and inhibits expression of p70 S6 kinase, useful for treating a condition associated with p70 S6 kinase, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Optionally 2'-methoxyethyl (2'-MOE) nucleotides.
All 2' MOE cytosines are 5-methylcytosine"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note= "Optionally 2'-methoxyethyl (2'-MOE) nucleotides.
ll 2' MOE cytosines are 5-methylcytosine"
                                                                                                                                                                                                  Human, p70 S6 kinase; SK6; p70/p85 s6 kinase; pp70s6k;
p70/p85 ribosomal S6 kinase; serine-threonine kinase;
ribosomal S6 protein phosphorylation; protein synthesis;
cell cycle progression; immune response; signalling cascade;
cancer progression; lipotoxic disorder; obseity; metabolic disorder;
hyperproliferative disorder; cancer; cyrostatic; expression inhibition;
phosphorothioate; antisense oligonucleotide; ss.
              Gaps
                                                                                                                                                                                Human p70 S6 kinase phosphorothioate antisense oligo, SEQ ID NO:21.
              .;
0
  Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                  GACATGTGGGGTGTGGGCTG 1172
                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= b
mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                     OTHER
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/mod_base= OTHER
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                                                      dacarcredrererere
                                                                                                                ВР
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80.08;
                                                                                                               ABZ74963 standard; DNA; 20
                                                                                                                                                          (first entry)
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 Best Local Similarity 80.0
Matches 16; Conservative
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*tag=
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
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                                                                                                                                                          10-MAY-2003
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protein, which in turn stimulates protein synthesis. p70 S6 kinase function is essential for cell cycle progression, and has also been implicated in the regulation of the immune response. p70 S6 kinase is itself activated via phosphorylation, a process influenced by upstream signalling cascades and by hyperinsulinaemia, and may play a role in the progression of colon cancer and in the development of lipotoxic disorders and obesity. The oligonoclecides of the invention are useful for the prevention and treatment of conditions associated with p70 S6 kinase, such as hyperproliferative disorders such as cancer, and metabolic disorders. They are also useful in research and diagnostics for modulating the expression of p70 S6 kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention comprises DNA sequences that show enhanced expression in human neuroblastroma with good prognosis. The DNA sequences of the invention are useful in clarifying good/poor prognosis of neuroblastoma. The present DNA sequence was used in the exemplification of the invention
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80.0%; Pred. No. 9.3e+02;
iive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.38+02;
tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 11 C; 3 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neuroblastoma; prognosis; ds; oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neuroblastoma-related DNA sequence #183.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     561 CCGCCGCCTCCGTCGTCA 580
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HISAMITSU PHARM CO LTD.
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24-AUG-2001; 2001JP-00255260.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-MAY-2002; 2002WO-JP005295.
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Best Local Similarity 80.0
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 80.0
nes 16, Conservative
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ABQ80265 standard; cDNA; 20
       (first entry)
          #2
          FLT-4 primer
        27-JUN-2003
     ABQ80265;
RESULT 1401
ABQ80265/c
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PCR; nervous system; platelet-derived growth factor; PDGF; psychosis; vascular endothelial growth factor; VBGF; neural; stem cell; memory; progenitor cell; neurodegeneration; ischaemia; neurological trauma; neuropsychiatry; learning; Parkinson's disease; Huntington's disease; Amyotrophic Lateral Sclerosis; spinal ischaemia; lschaemic stroke; spinal cord injury; cancer-related; schizophrenia; Alzheimer's disease; depression; anxiety; phobia; stress; cognitive function; aggression; anxiety; phobia; stress; cognitive function; aggression; multi-infarct; dementia; Lewy body; age related; geriatric; growth; epilepsy; brain injury; multiple sclerosis; autism; differentiation; attention deficit disorder; narcolepsy; amplify; ss.

Homo sapiens

WO2003024478-A1.

27-MAR-2003.

19-SEP-2002; 2002WO-IB003998

19-SEP-2001; 2001US-0323381P. 28-SEP-2001; 2001US-0326044P.

(NEUR-) NEURONOVA AB

Plate K, Schanzer A, Wachs F; Kuhn GH. Delfani K, Janson AM, Zhao M;

WPI; 2003-354563/33

Use of platelet-derived growth factor, vascular endothelial growth factor, or their modulators for modulating neural stem cell or neural progenitor cell activity, particularly for treating e.g. Alzheimer's, ischemia or stroke.

Example 11; Page 77; 119pp; English.

The sequences given in ABO80256-69 are primers which were used to identify the presence of vascular endothelial growth factor (VEGF), or the VEGF receptors, FIK1, FIT-1 and FIT-4 in human stem cells (BEC). The method of the invention for alleviating or reducing a symptom of a disease or disorder of the nervous system comprises administering prowth factor (VEGF), a combination of PDGF and VEGF, or a PDGF or VEGF agonist, to a patient in order to modulate neural stem cell or neural progenitor cell activity in vivo. The method is useful for alleviating or reducing the symptoms of a disease or disorder of the nervous system, e.g. neural progenitor disorders, neural stem cell disorders, neural progenitor disorders, neural stem cell disorders and enemory affective disorders, neuropsychiatric disorders or learning and memory affective disorders, neuropsychiatric disorders or learning and memory affective disorders, spinal cord injury, or cancer-related brain/ spinal cord injury, schizophrenia and other psychoses, depression, bipolar chancer, anxiety syndromes/disorders, appression, bipolar chancer, borderine personality disorder, cerebral palsy, life style disorder, borderline personality disorder, cerebral palsy, life style drug, multi-infarct dementia, Lewy body dementia, age related/geriatric

10-OCT-2001; 2001US-00973827. 07-OCT-2002; 2002WO-US032181.

WO2003030617-A2

17-APR-2003.

(ISIS-) ISIS PHARM INC

Cowsert LM;

Monia BP,

WPI; 2003-381663/36.

New antisense oligonucleotides for modulating CREB (CAMP response element binding protein) gene expression, useful for preventing or treating e.g. cancers, a disease arising from aberrant apoptosis, or neuronal

Claim 3; Page 74; 91pp; English

disorders.

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dementia, epilepsy and injury related to epilepsy, spinal cord injury, brain arelated brain/spinal cord injury, anti-cancer treatment related brain/spinal cord injury, infection and inflammation related brain/spinal cord injury, environmental toxin related brain/spinal cord injury, environmental toxin related brain/spinal cord injury, environmental toxin related brain/spinal cord injury, multiple sclerosis, autism, attention deficit disorders, narcolepsy or sleep disorders. The PDGF and/or VBGF, is useful in the manufacture of a medicament for alleviating or treating these diseases or disorders, accelerating growth of neural stem cells or neural progenitor cells, or inducing proliferation or differentiation of these cells. This primer gives an estimated band size of 378 bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; CREB; cAMP response element binding protein; CREB1; bZIP; basic leucine zipper; transcription factor; intracellular signalling; spermatogenesis; circadian rhythm; memory; apoptosis; hyperproliferative disorder; cancer; tumour; blood; soft tissue; apoptosis related disease; neuronal disorder; chromosome 2q32.3-34; cytostatic; neuroprotective; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and 3' ends,
                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "This oligonucleotide has a phosphorothioate backbone with all cytosine residues being 5-methylcytosines. Optionally, it also has 2-methylcytosines. Optionally, it also has 2-wethyoxyethyl (2'-MOE) wings at the 5' and 3' ends which are 5 nucleotides in length."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human CREB phosphorothioate antisense oligonucleotide, SEQ ID NO:29.
                                                                                                                                                                                                                                                                           ;
0
                                                                                                                                                                                                                                      0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
ative 0; Mismatches 4; Indele
                                                                                                                                                                                                     Sequence 20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                             514 CTGGAGAGCTGACCCTCAA 533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                          20 CIGGIGAAGCIGCCCGIGAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                        ACF33771 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-SEP-2003 (first entry)
                                                                                                                                                                                                                                                                             16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACF33771;
                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1402
                                                                                                                                                                                                                                                                             Matches
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ID AC
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The invention relates to antisense oligonucleotides (ACF33752-ACF33779)
targeted to the human CREB (CAMP response element binding protein) gene,
which inhibit its expression. The oligonucleotides were designed to
target different regions of human CREB mRNA, and were analysed for their
effect on CREB expression by quantitative real-time PCR. CREB (also known
as CREB1) is a member of the basic leucine zipper (bZIP) family of
transcription factors, and activates transcription of target genes in
response to a diverse array of stimuli including peptide hormones, growth
actors and protein kinases. It is a component of intracellular
signalling events which regulate a wide variety of biological functions,
including spermatogenesis. Circadian rhythms and memory. Overexpression
of CREB has been found to induce apoptosis in certain cells, although
cresponse to drug dependency, as it has been found to mediate morphine-
induced upregulation of the CAMP pathways that contribute to opiate
development of drug dependency, as it has been found to mediate morphine-
induced upregulation of the cAMP pathways that contribute to opiate
dependency. The oligonucleotides of the invention are useful for
and soft tissuely, diseases or conditions arising from aberrant apoptosis,
or neuronal disorders (particularly cancer, e.g., those of blood
and soft tissuely, diseases or conditions arising from aberrant apoptosis,
or neuronal disorders The present sequence represents a human H-ras
phosphorothioate antisense oligonucleotide used as a positive control in
incention of the concentration for a particular cell
Mon May
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Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;

Gaps . 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ive 0; Mismatches 4; Indels Query Match Best Local Similarity 80.0 Matches 16; Conservative

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294 TICTGCACGGGCCCCACTCA 313 Trarecarecececeaca 20 Н

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ABT32380 standard; DNA; 20 BP 08-MAY-2003 ABT32380; RESULT 1403 ABT32380

(first entry)

Neuroblastoma-related oligonucleotide #157.

Neuroblastoma; prognosis; spontaneous regression; primer; probe; ds; high malignancy

WO200297093-A1.

05-DEC-2002

30-MAY-2001; 2001JP-00162775. 24-AUG-2001; 2001JP-00255226. 30-MAY-2002; 2002WO-JP005294.

(CHIB-) CHIBA PREFECTURE. (HISM) HISAMITSU PHARM CO LTD.

Nakagawara A;

WPI; 2003-140476/13.

Nucleic acids having higher expression in human neuroblastoma with poor prognosis for diagnostic prediction of neuroblastoma prognosis.

Example 5; Page 27; 111pp; Japanese.

The invention comprises nucleic acids that show increased expression in

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human neuroblastomas with poor prognosis over those with a good prognosis. The nucleic acids of the invention are useful as a tool for distinguishing neuroblastomas with a favourable prognosis (spontaneous regression) from neuroblastomas with a poor prognosis (high malignancy). The DNA sequences ABT32224 - ABT32571 represent oligonucleotides used in an example of the invention
                                                                                                                                                                                                                                                                                                                                                                                                         BCL2-associated X, BAX; nootropic, neuroprotective; antiparkinsonian, anticonvulsant; ophthalmological; antidiabetic; virucide; antisense therapy; BAX antagonist; BAX inhibitor; familial amylotrophic lateral sclerosis; Alzheimer's disease; PaxKinson's disease; hodgkin's disease; cartilage-hair hyperplasis; adiabetes-associated ocular disorder; scrapie infection; aberrant apoptosis; human; phosphorothioate; ss.
                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "phosphorothioate linkages, and all cytidine
                                                                                                                                                                                                                                                                                                                                                                               Human BAX chimeric phosphorothicate oligonucleotide SEQ ID NO:27.
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                                                                                                                                 0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
                                                                                                      Sequence 20 BP; 7 A; 7 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        residues are 5-methylcytidines
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
16. .20
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/note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                          316 TCTGCACCAGAGATTGTGCA 335
                                                                                                                                                                                                              1 TCTGCACCAGAGAATCCACA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER
                                                                                                                                                                                                                                                                                             ADA20854 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JUL-2001; 2001US-00908147.
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                                                                                                                  Query Match
Best Local Similarity 80.00,
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*tag=
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modified_base
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                                                                                                                                                                                                                                                                                                                        ADA20854;
                                                                                                                                                                                                                                                                1404
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ADA20854/
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Example 15; Page 85; 139pp; English.
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The present invention describes a compound (1) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding BCL2-associated X (BAX) protein, where the compound specifically hybridises with the mucleic acid molecule encoding BAX protein. The compound specifically hybridises with at least 8-nucleobase portion of an active site on a nucleic acid molecule encoding BAX protein. Also described: (1) a composition comprising (1) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of BAX protein in cells or tissues comprising contacting the cells or tissues with (1); and (3) treating an animal having a disease or condition associated with BAX protein comprising administering to the animal (1) so that expression of BAX protein is inhibited. (1) has nootropic, neuroprotective, antiparxinsonian, anticonvulsant, ophthalmological, antidiabetic and virtugarishing and en be used in antisense therapy, and as a BAX antiagonist. The antisense compounds (1) are useful for modulating the expression of BAX protein, and for treating a disease or condition associated with BAX protein, and for treating a disease or condition associated with BAX protein, and for treating a disease or condition associated with BAX protein, e.g. familial amylotrophic lateral schedular hyperplasia, diabetes-associated ocular disorders or scrapie infection, or a condition that arises from aberrant apoptosis. The present sequence represents a human BAX chimmeric phosphorothhoate present invention.
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Sequence 20 BP; 2 A; 8 C; 7 G; 3 T; 0 U; 0 Other;

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Gaps
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Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
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ADA20960 standard; DNA; 20 BP ADA20960; 1405 DA20960 RESULT

(first entry) 20-NOV-2003

Mouse BAX chimeric phosphorothioate oligonucleotide SEQ ID NO:133.

BCL2-associated X; BAX; nootropic; neuroprotective; antiparkinsonian, anticonvulsant; ophthalmological; antidiabetic; virucide; antisense therapy; BAX antisense therapy; BAX inhibitor; familial amylotrophic lateral sclerosis; Alzheimer's disease; Parkinson's disease; Hodgkin's disease; cartilage-hair hyperplasia; diabetes-associated ocular disorder; scrapie infection; aberrant apoptosis; mouse; phosphorothioate; ss.

Synthetic

Mus musculus

/note= "phosphorothioate linkages, and all cytidine residues are 5-methylcytidines" Location/Qualifiers /*tag= b /mod_base= OTHER ...5 '*tag= a Key modified_base nodified base

/mod_base= OTHER /note= "2'-0-methoxyethyls" 16. .20 /*tag= c /mod_base= CTHER /note= "2'-0-methoxyethyls" modified_base

WO2003008543-A2

13-JUL-2002; 2002WO-US022417.

17-JUL-2001; 2001US-00908147.

(ISIS-) ISIS PHARM INC

Zhang H, Watt AT;

New antisense compounds, useful for modulating the expression of BCL2-associated X (BAX) protein or for treating a disease or condition associated with BAX protein, e.g. Parkinson's disease, Hodgkin's disease WPI; 2003-239321/23

Example 17; Page 94; 139pp; English.

or Alzheimer's disease

The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding BCI2-associated X (BXX) protein, where the compound specifically hybridises with the nucleobase portion of an active site on a nucleic acid molecule encoding BAX protein. The compound specifically hybridises with at least 8-nucleobase portion of an active site on a nucleic acid molecule encoding BAX protein. Also described: (1) a composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of BAX protein in cells or tissues comprising contacting the cells or tissues with (I), and (3) treating an animal having a disease or condition associated with BAX protein in comprising administering to the animal (I) so that expression of BAX protein is inhibited. (1) has nootropic, neuroprotective, antiparkinsonian, anticonvilsant, ophthalmological, antidiabetic and virtida antisense compounds (I) are useful for modulating the expression of BAX protein, and for treating a disease or condition associated with BAX protein, and for treating a disease or condition associated with BAX protein, and for treating a disease or condition associated with BAX protein, e.g. familial amylotrophic lateral cartiage-hair hyperplasia, diabetes-associated coular disorders or scraple infection, or a condition that arises from aberrant apoptosis. The present sequence represents a mouse BAX chimeric phosphorothioate present invention.

Seguence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps 0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.38+02; tive 0; Mismatches 4; Indels Query Match
Best Local Similarity 80.0
Matches 16; Conservative

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RESULT 1406 ACF39671

ACF39671 standard; DNA; 20 BP.

ACF39671;

(first entry) 29-SEP-2003

MHC class II transactivator antisense oligonucleotide SEQ ID NO:74.

Human, major histocompatibility complex class II transactivator;
MHC class II transactivator; antisense modulation; immunosuppressive;
antimicrobial; antidiabetic; antirheumatic; antiarthritic; cytostatic;
nootropic; neuroprotective; immunostimulant; autoimmune disorder;
MHC Class II transactivator inhibitor; infection; transplant rejection;
diabetes; rheumatoid arthritis; cancer; Alzheimer's disease;
multiple sclerosis; severe combined immunodeficiency disease;

AAL61864 standard; DNA; 20 BP.

RESULT 1407 **AAL**61864 AAL61864;

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The present invention describes a compound (I) that is 8-50 nucleobases in length: (a) targets a nucleic acid molecule encoding major histocompatibility complex (MHC) class II transactivator, and inhibits the expression of MHC class II transactivator; or (b) specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding MHC class II transactivator. (I) has immunosuppressive, antimicrobial, mutilabetic, antirheumatic, antiarthritic, cytostatic, nootropic, neuroprotective and immunostimulan activities, and can be used as an MHC class II transactivator and immunostimulan activities, and can be used as an utilitiesnee oligonuclectides can be used for treating an animal having a disease or condition associated with MHC class II transactivator, e.g. autoimmune disorder or infection. The antisense oligonuclectides can be used for inhibiting the expression of MHC class II transactivator in cells or tissues. In particular, these diseases include transplant celts or inhibiting the expression of MHC class II transactivator in cells or tissues. In particular, these diseases include transplant multiple solerosis, or severe combined immunodeficiency disease. The antisense compounds are useful for diagnostics, prophylaxis, or as antisense compounds are useful for diagnostics, prophylaxis, or as antisense compounds are useful for diagnostics prophylaxis, or as creared reagents or kies. The present sequence represents a human MHC class II transactivator chimmeric phosphorothioate antisense invention or which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides for modulating MHC class II transactivator gene expression, particularly useful for treating autoimmune disorders such as transplant rejection, Alzheimer's disease, or multiple sclerosis,
                                                                                                                                            /*rag= a/mod_bos= OTHER
/mod_bos= OTHER
/nore= "phosphorothioate linkages; all cytidine residues
are 5-methylcytidines"
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  phosphorothioate; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                          /*tag= c
/mod_base= OTHER
/note= "2'-0-methoxyethyls"
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/note= "2'-O-methoxyethyls"
16. .20
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                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-DEC-2001; 2001US-00006366.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-DEC-2002; 2002WO-US038616.
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                                                                                                                      *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-577294/54.
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                                                                                                                 modified_base
                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                        modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             or infection.
                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bennett FC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JUN-2003
                                                              Synthetic
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The invention relates to antisense compounds targetted to the nucleic acid encoding the G protein-coupled receptor FTBR-LD-2 (endothelin type b receptor-like protein-2) to inhibit its expression. FTBR-LD-2 is also known as endothelin-binding receptor-like protein-2, FTBR-LIke protein 2 and G-protein coupled receptor 37 like 1 (FBR3/LI). Antisense compounds of the invention are useful for treating hyperproliferative disorders (especially cancer) and cardiovascular diseases especially angiogenesis, arthroposlerosis, hypertension, cerebral vascular disease, stroke and acute proliferative nephropathy. The present sequence is an antisense oligonucleotide targetted to human ETBR-LD-2 DNA
                                                                                                                 Human, G protein-coupled receptor; hyperproliferative disorder; GPR37L1; endothelin type b receptor-like protein-2; cerebral vascular disease; antisense; endothelin-binding receptor-like protein-2; atherosclerosis; cardiovascular disease; ETR-LP-2; G-protein coupled receptor 37 like 1, acute proliferative nephropathy; ETBR-like protein 2; cancer; stroke;
                                                                                                                                                                                                                                                                                                                                                                       /note= "Phosphorothioate backbone; All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotides which bind the nucleic acid encoding the G protein coupled receptor BTBR-LP-2 (endothelin type b receptor-like protein-2 receptor), useful for treating e.g. cancer and cardiovascular diseases.
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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.8%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 9.3e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                  Human ETBR-LP-2 antisense oligonucleotide ISIS #204290.
                                                                                                                                                                                                             angiogenesis; hypertension; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 20 BP; 7 A; 7 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Page 80; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                      mod_base= OTHER
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                                              22-SEP-2003 (first entry)
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                                                                                                                                                                                                                                                Homo sapiens.
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0; Gaps

0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels

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Best Local Similarity 80.0 Matches 16; Conservative

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The invention relates to antisense compounds targetted to the nucleic acid encoding the G protein-coupled receptor ETBR-LP-2 (endothelin type b receptor-like protein-2) to inhibit its expression. ETBR-LP-2 is also known as endothelin-binding receptor-like protein-2, ETBR-LP-2 is also and G-protein coupled receptor 37 like i (GR3711). Antisense compounds of the invention are useful for treating hyperproliferative disorders (especially cancer) and cardiovascular diseases especially angiogenesis, atherosclerosis, hypertension, cerebral vascular disease, stroke and acute proliferative nephropathy. The present sequence is an antisense oligonucleotide targetted to human ETBR-LP-2 DNA
                                                                                                                                                                                                                                                  Human, G protein-coupled receptor; hyperproliferative disorder; GPR37L1; endothelin type b receptor-like protein-2; cerebral vascular disease; antisense; endothelin-binding receptor-like protein-2; atherosclerosis; cardiovascular disease, ETBR-LP-2; G-protein coupled receptor 37 like 1; acute proliferative nephropathy; ETBR-like protein 2; cancer; stroke; anglogenesis; hypertension; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Phosphorothioate backbone; All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New oligonucleotides which bind the nucleic acid encoding the G protein coupled receptor ETBR-LP-2 (endothelin type b receptor-like protein-2 receptor), useful for treating e.g. cancer and cardiovascular diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                       Human ETBR-LP-2 antisense oligonucleotide ISIS #204289.
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                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
1...20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 15; Page 80; 106pp; English
1049 GAGCCAAGTCAATCCCAACA 1068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= c
/mod_base= OTHER
                              GAACCAAGTCCATCCCTAGA 20
                                                                                                                    BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-DEC-2002; 2002WO-US038520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-DEC-2001; 2001US-00003126
                                                                                                                    AAL61863 standard; DNA; 20
                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    *tag= p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Freier SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-558997/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003050244-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                              modified base
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                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                      22-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                     AAL61863;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic context dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                   Gaps
                                                                                                                                                                                                                                                                                                             Immunostimulatory, antiinflammatory, dermatological, antipsoriatic, antiulcer; gene therapy, vaccine; non-allergic inflammatory disease, psoriasis; eczema; allergic contact dermatitis; latex dermatitis; laflammatory bowel disease, ulcerative colitis; Crohn's disease; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gabe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treating non-allergic inflammatory diseases, such as psoriasis, eczallergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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Pred. No. 9.3e+02;
0; Mismatches 4; Indels
 Length 20;
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0.8%; Score 13.6; DB 1;
80.0%; Pred. No. 9.3e+02;
cive 0; Mismatches 4;
                                                                                                                                                                                                                                                                               Immunostimulatory nucleic acid #235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 15; 229pp; English.
                                                               1052 CCAAGTCAATCCCAACAAAG 1071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   555 CCTCAGCCGCCGCCTCCGTC 574
                                                                                               1 ccaagrcarccragacae 20
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                                                                                                                                                                                ACD99549 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-MAR-2002; 2002US-00112653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAR-2001; 2001US-0279642P
                                                                                                                                                                                                                                                25-SEP-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 80.05
Matches 16; Conservative
Query Match
Best Local Similarity 80.0
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Krieg AM, Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (KRIE/) KRIEG A M. (BERG/) BERG D J.
                                                                                                                                                                                                                                                                                                                                                                                                                                  US2003050268-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                  ACD99549;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA15368
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Mouse; PCR; primer; ss; HYPLIP1; FCHL1; variation; lipid disorder; allee; anti-lipid disorder; anti-cancer therapy; gene therapy; familial combined hyperlipidaemia; coronary artery disease; atherogenic lipoprotein phenotype; hyperapobetalipoproteinaemia; hypertriglyceridaemia; low density lipoprotein subclass B; LDL; familial dyslipidaemic hypertension; syndrome X; hypercholesterolaemia; obesity; insulin resticance; cancer; cytostatic; antilipaemic; hypotensive; anorectic.
                                                                                                    Mouse HYPLIP1 locus PCR primer #308.
06-NOV-2003 (first entry)
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Mus sp.

03-APR-2003.

07-SEP-2001; 2001US-00949428

22-JUN-2000; 2000US-0213322P.

(BODN/) BODNAR J S. (CAST/) CASTELLANI L W. (CHAT/) CHATTERJEE A. (JONG/) JONG P D. (LUSI/) LUSIS A J. (OHNE/) OHNEN J. (ROSK), ROSS D. (TAFU/) TAFURI S. (WUCC/) WU C.

Chatterjee A, Jong PD, Lusis AJ; , Wu C; Bodnar JS, Castellani LW, C Ohmen J, Ross D, Tafuri S,

WPI; 2003-540780/51

Novel isolated polynucleotide comprising a mouse or human familial combined hyperlipidemia 1 gene having a variation that is associated with a lipid disorder, useful for identifying susceptibility to the lipid

Claim 11; Page 40; 63pp; English

The invention discloses isolated polymucleotides comprising mouse HYPLIPI control as sequence, mouse HYPLIPI genomic DNA, or the homologous human channed mouse HYPLIPI genomic DNA, or the homologous human control and the sequence is associated with a lipid disorder. Also claimed is an isolated polyperlipidaemia 1 (FCHLI) game, where a variation in the sequence, or a variant form of a fully defined human FCHLI amino acid sequence, where the variant form of the mouse HYPLIPI amino acid sequence, where the variant is associated with the lipid disorder, an isolated polymucleotide having at least 12 contiguous nucleotides of the variation position, an isolated polypeptide comprising 4 contiguous amino acids of the encode polypeptides, where the 4 contiguous amino acids of the encode polypeptides, where the 4 contiguous amino acids span the variation position, a kit for the detection of the FCHLI locus comprising, an isolated anilody, identifying susceptibility to a lipid disorder which comprises comparing the nucleotide sequence of the difference between the suspected allele and the wild-type sequence identifies a sequence variation of FCHLI nucleotide sequence and a pharmaceutical composition. Also disclosed is a transgenic animal which carries an altered HYPLIPI or FCHLI allele and a method for screening drugs for inhibition or restoration of FCHLI gene function as an antilipid disorder associated with expression of FCHLI, for diagnosis or prognosis of predisposition to lipid disorder, and cancer and for prognesse of predisposition to lipid disorder, and cancer and for treating a lipid disorder such as familial combined hyperlipidaemia, occonary artery disease, atherogenic lipoporteinaemia, low density

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lipoprotein (LDL) subclass B, familial dyslipidemic hypertension, syndrome X, hypercholesterolaemia, obesity, insulin resistance and cancer. The sequence presented is a PCR primer which was used to amplify part of the mouse HYPLIPI locus.
                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Embryonic stem cell; ES cell; mouse; differentiation; nerve or
panoreatic islet cell; cell transplant therapy; antidiabetic;
neuroprotective; nootropic; PCR; primer; ss.
                                                                                                                                                                                                                 Score 13.6; DB 1; Length 20; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                           4; Indels
                                                                                                                                                      Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Murine embryonic cell line Ptc PCR primer #1.
                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (OKUM-) OKUMA CONTACTLENS KENKYUSHO YG. (INOU/) INOUE K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inoue K, Kim D, Gu Y, Ishii M;
                                                                                                                                                                                                                                                                                                                                         16 GGATGGACAGGAATGCAGAG 35
                                                                                                                                                                                                                                                                                                                                                                                          20 GGATGGAGAGGCATCCTGAG 1
                                                                                                                                                                                                                    0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-JAN-2002; 2002US-00054789.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-JAN-2003; 2003WO-JP000699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACF04246 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-NOV-2003 (first entry)
                                                                                                                                                                                                                 Query Match
Best Local Similarity 80.03
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-598750/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003062405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-JUL-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACF04246;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1411
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The present invention relates to a method of inducing differentiation of mammalian embryonic stem cells into functioning cells, which comprises culturing embryonic stem cells in a medium comprisely which comprises culturing embryonic stem cells in a medium comprise, the invention cator and basic fibroblast growth factor. In particular, the invention cells are to the differentiation of murine embryonic stem cells. The method is useful for inducing differentiation of mammalian embryonic stem cells. The method cells into functioning cells. Other methods are useful for treating a mammalian patient having disorders in pancreatic function, and in nerve function. The cells are pancreatic slet like cell clusters and nerve like cells. Functioning cells induced from embryonic stem cells using the present method may be used for treating disorders in pancreatic islet function creation disorders or disease), neuronal degeneration (e.g. Alzheimer's disease) cells are useful not only for cell transplant therapy, but for in vitro creation, and for safety evaluation of new drugs. The present sequence is a PCR primer used in the exemplification of the invention

Inducing differentiation of mammalian embryonic stem (ES) cells into functioning cells, for treating e.g. diabetes, comprises culturing ES cells in a medium containing leukemia inhibitor factor and basic fibroblast growth factor.

Example 5; Page 67; 70pp; English.

BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other; Sequence 20 ö

Gaps

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4; Indels

Pred. No. 9.3e+02; 0; Mismatches

80.08;

16; Conservative

Best Local Similarity Matches 16; Conserv

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The invention discloses a method for amplifying expressed genetic sequences from genomic DNA (gDNA) from mammalian or higher order plant species. The method involves identifying a 3 untranslated region (UTR) of a gDNA sequence, designing probe, performing a second PCR to amplify product by size differentiation and performing a second PCR to amplify the predetermined sequence. Also claimed is a biological analysis device, comprising a substrate and an array of a set of expressed genetic sequences, located on the substrate, which are generated by the method are deposited an array of biosites of genomic DNA fragments having the sequence of at least one exon, and absent polyadenine and vector sequences to exon, and absent polyadenine and vector sequences where the genomic DNA fragments have a sequence length of from about 75-2000 nucleotides. The method is efficient for amplifying gene sequences, eaches a large-scale production of gDNA sequences, enables large-scale production of gDNA secuences is generated by including in microarray formats, fabricates high-density DNA arrays of printing in microarray formats, fabricates high-density DNA arrays of enhanced, watchy varying genetic content and abstains from using RNA-derived sequences by simple PCR amplifications without cloning. The method produces amplified sequences that have greater specificity and size consistency than that observed with cDNA fragments, and allows for is a Type II gene specific primer.
                                                                                                                                                                                                                                                                                                                                                                                    PCR; primer; ss; genomic DNA; gDNA; untranslated region; UTR; DNA high-density microarray; biosite; large scale production; gDNA probe; microarray; Type II primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Amplifying expressed genetic sequences from genomic DNA of mammalian or higher order plant species for printing on DNA microarrays, involves using the 3' untranslated region of the gene sequence.
                                               Gaps
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0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                  Sense PCR primer used to amplify USP15.
                                                                               614 CCTACATTAAGCTGGACAAA 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 8; 15pp; English
                                                                                                             1 CCTCCTTTACGGTGGACAAA 20
                                                                                                                                                                                                            .607/c
ACH66607 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-OCT-2001; 2001US-00972469.
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                                                                                                                                                                                                                                                                                                         (first entry)
Query Match
Best Local Similarity 80.0°
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JS2003073085-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (LAIF/) LAI F.
(ZHOU/) ZHOU D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lai F, Zhou D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                         16-0CT-2003
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                                                                                                                                                                                                                                                                  ACH66607;
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The invention describes an isolated polypeptide (I) comprising a variant form d a mouse HYPLIPI polypeptide sequence (S1) or a human FCHLI polypeptide sequence (S1) or a human FCHLI polypeptide sequence (S1) or a human FCHLI polypeptide sequence (S1) or diagram or a read or a sequence having at least 65 % sequence identify to (S1) or (S2). A composition comprising the expression of FCHLI. FCHLI gene or HYPLIRI gene and its product are useful for the study of metabolic pathway and cellular mechanism to identify other genes, receptors and relationships that contribute to lipid disorder and cancer. FCHLI gene or its fragments are useful in gene the ready to increase the amount of the expression products of the gene for the treatment of lipid disorder or cancerous cells. The sequence variation of FCHLI gene or HYPLIPI gene is also useful in the disgnosis and prognosis of predisposition to lipid disorder and cancer. Antisense collymuclectide sequences are useful in preventing or diminishing the expression of HYPLIPI or mouse HYPLIPI gene.
                                                                                                                                                                                                                               cytostatic, antilipemic; gene therapy; peptide therapy; HYPLIPI; FCHLI; cancer; metabolic pathway; cellular mechanism; lipid disorder; familial combined hyperlipidaemia; mouse; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human FCHLI or mouse HYPLIPI polypeptide, useful for drug
screening, peptide therapy of lipid disorder or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lusis AJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jong PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chatterjee A,
Wu C;
742 ACCGCCATCCGGGAAGTGTC 761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 11; Page 38; 56pp; English.
                          20 ACCACCAGCAGGAAAGTGTC 1
                                                                                            930/c
ADB95930 standard; DNA; 20 BP.
                                                                                                                                                                                                 Mouse HYPLIP1 PCR primer #308.
                                                                                                                                                                                                                                                                                                                                                                                     07-SEP-2001; 2001US-00949427.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Castellani LW, C'
oss D, Tafuri S,
                                                                                                                                                                                                                                                                                                                                                                                                                    08-SEP-2000; 2000US-0231322P.
                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                               BODNAR J S.
CASTELLANI L W.
CHATTERJEE A.
JONG P D.
LUSIS A J.
OHMEN J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ross D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ROSS D.
TAFURI S.
                                                                                                                                                                                                                                                                                                                          US2003054418-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (OHME/) OHMEN
(ROSS/) ROSS I
(TAFU/) TAFURI
(WUCC/) WU C.
                                                                                                                                                                 04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                         20-MAR-2003.
                                                                                                                                     ADB95930;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BODN/)
(CAST/)
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                                                                                                                                                                                                                                                                                             Mus sp.
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0.8%; Score 13.6; DB 1; Length 20;

Query Match

0.8%; Score 13.6; DB 1; Length 20;

Query Match

Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;

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Example 8; Page 129; 222pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   154 CIGICAAIGACACICCGAGG 173
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                                                                                                                                                          (HELI-) HELIX RES INST.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                       28-MAR-2002; 2002EP-00007401.
                                                                                                                            05-NOV-2001; 2001JP-00379298.
25-JAN-2002; 2002US-00350978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC65807 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 European Patent Office.
                                                                                                                                                                                         Sugiyama T,
                                                                                                                                                                                                   Yamamoto J, Isono Y,
Seki N, Yoshikawa T,
                                                                                                                                                                                                                                    WPI; 2003-450961/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                              EP1308459-A2.
                                          Homo sapiens
                                                                                 07-MAY-2003.
                                                                                                                                                                                         Isogai T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC65807;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1416
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  셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-3 nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnoshing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                  Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
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            Gaps
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           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical; diagnostic; gene therapy; tissue regeneration;
                                                                                                                                                                                         ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.6; DB 1; Length 20;
llarity 80.0%; Pred. No. 9.3e+02;
Conservative 0; Mismatches 4; Indels
Similarity 80.0%; Fred. No. 9.3e+02;
16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 6 C; 14 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                            Fouron Y;
                                                                                                                                                                      Immunostimulatory nucleic acid #232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 8; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sss cereacecececerees s74
                              16 GGATGGACAGGAATGCAGAG 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ccecceccecceccecce 1
                                             GGATGGAGAGGCATCCTGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Clone specific PCR primer #136.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bb.
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                                                                                                                                                                                                                                                                                                            03-FEB-2000; 2000US-0179991P.
                                                                                                                                                                                                                                                                                       02-FEB-2001; 2001US-00776479.
                                                                                                                                                                                                                                                                                                                                                                           Bratzler RL, Petersen DM,
                                                                                                       ADB36618 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB65935 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                              (BETE/) BRATZLER R L. (PETE/) PETERSEN D M. (FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-657977/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
les 16; Conserv
 Best Local Similarity
                                                                                                                                                                                                                                              US2003087848-A1
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                                                                                                                                                                                                                                                                   38-MAY-2003.
                                                                                                                                                                                                                          Synthetic.
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Best Local S:
Matches 16
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                                                                                                                            ADB36618;
                                                   20
                                                                                 RESULT 1414
          Matches
                                                                                              ADB36618/
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The invention discloses a polynucleotide comprising a sequence selected from 1970 fully defined nucleotide sequences which encode novel polypeptides. Also claimed is a polypeptide encoded by the polynucleotide or its partial peptide, an antibody binding to the polypeptide or peptide of the polynucleotide, immunologically assaying the polypeptide or peptide with the antibody of the encoded by contexting the polypeptide or peptide with the antibody of the encoded by contexting the polypeptide or peptide with the antibody of the encoded by contexting the polynucleotide in an expressible manner and an antisense polynucleotide, or as a probe for detecting the polynucleotide. The polynucleotide in an encoded is useful as a primer for synthesisting the polynucleotide and encoded for detecting the polynucleotide. The polynucleotide and encoded for detecting the polynucleotide. The polynucleotide and encoded for detecting the polynucleotide. The polynucleotide and encoded genes may be included in them, for developing a diagnostic marker or medicines for regulation of their expression and activity, or as targets of gene thrapy. The genes are involved in tissue and/or cell regeneration. Membrane proteins, disease-related proteins are involved in tissue and/or cell cranscription-related proteins, disease-related proteins and genes encoding them can be used as indicators for diseases (e.g. osteoporosis, reanscription-related proteins, disease-related proteins and genes concing them can be used as indicators for diseases (e.g. osteoporosis, neurological diseases, cancer, tumours. The cDNA may be used to regulate the activity or expression of the encoded protein to treat diseases. The sequence presented is clone specific PCR primer which was used in the sequence data for this patent is not represented in the printed sequence data for this patent is not represented in the printed or proteins and proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Otsuki T, Wakamatsu A, Sato H, Ishii S;
Hio Y, Otsuka K, Nagai K, Irie R, Tamechika I;
Otsuka M, Nagahari K, Masuho Y;
cell regeneration, membrane protein; signal transduction-related protein;
transcription-related protein; osteoporosis; neurological disease;
cancer; tumour; primer; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polynuclectides and polypeptides, useful for developing a diagnostic marker or medicines for regulation of their expression and activity, or as targets of gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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80.0%; Pred. No. 9.3e+02;
iive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 7 A; 4 C; 7 G; 2 T; 0 U; 0 Other;
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The invention comprises antisense oligonucleotides that are targeted to the nucleic acid encoding transforming growth factor beta (TGF-beta) receptor II. The antisense oligonucleotides of the invention are useful for treating: hyperproliferative disorders (e.g. breast cancer), or an autoimmune disorder (e.g. prheumatoid arthritis). The present DNA sequence represents a 2.-O-methoxyethyl gapmer oligonucleotide with a phosphorothioate backbone that is targeted to mouse TGF-beta receptor II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             New compound having a sequence targeted to a nucleic acid encoding
Transforming growth factor beta-receptor II, useful for preparing a
composition for treating hyperproliferative disorder e.g., lung, liver,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; primer; cytostatic; antidiabetic; anorectic; cerebroprotective; neuroprotective; antiinflammatory; gene therapy; antisense therapy; thyromimetic; Novix; pathology; cancer; diabetes; obseity; endocrine disorder; CNS disorder; inflammatory disorder; chromosome mapping; tissue typing; predictive medicine.
                                                                mouse; antisense oligonucleotide;
transforming growth factor beta receptor II; TGF-beta receptor II;
hyperproliferative disorder; breast cancer; autoimmune disorder;
rheumatoid arthritis; 2'-0-methoxyethyl gapmer;
phosphorothioate backbone; ss; murine.
                                   Mouse TGF-beta receptor II targeted antisense oligonucleotide #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human NOVX polypeptide gene forward primer SEQ ID NO: 535.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 9 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; SEQ ID NO 103; 141pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         108 GCCCCCCCCATCCCCATGG 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         eccccccrccrccrcarae 20
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                                                                                                                                                                                                                                                                                 19-JUN-2002; 2002WO-US019665
                                                                                                                                                                                                                                                                                                                   21-JUN-2001; 2001US-00888361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC10516 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
18-DEC-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 80.0°
Matches 16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  colon or gastric cancer
                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                         Murray SF, Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-175279/17.
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                                                                                                                                                                                                             WO2003000656-A2
                                                                                                                                                                             Mus musculus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-DEC-2003
                                                                                                                                                                                                                                               03-JAN-2003
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ADC10516/c
BXXXXXXXXXXXXXXXX
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New isolated NOVX polypeptides and nucleic acid molecules useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, such as cancer, obesity, diabetes and inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Agee ML, Anderson DW, Berghs C, Casman SJ, Catterton E;
Dipippo VA, Edinger SR, Eisen A, Ellerman K, Gangolli EA;
Gerlach VL, Gorman L, Guo X, Harkmann JL, Hjalt T, Ji W, Kekuda R
Khramtsov NV, Li L, Liu X, Malyankar UM, Miller CE, Millet I;
Ort T, Padigaru M, Patturajan M, Pena CEA, Rastelli L, Rieger DK;
Rotheberg ME, Shenoy SG, Shimkets RA, Smithson G, Spaderna SK;
Spytek KA, Stone DJ, Vernet CAM, Zhong H, Zhong M, Alsobrook JP;
Burgess CE, Lepley DM;
                                    04-JUN-2001, 2001US-0295607P.
04-JUN-2001, 2001US-0295601P.
06-JUN-2001, 2001US-0295618P.
06-JUN-2001, 2001US-0295618P.
11-JUN-2001, 2001US-0295618P.
12-JUN-2001, 2001US-0295573P.
12-JUN-2001, 2001US-0295573P.
13-JUN-2001, 2001US-0295573P.
19-JUN-2001, 2001US-029958P.
19-JUN-2001, 2001US-029953P.
19-JUN-2001, 2001US-029959P.
22-JUN-2001, 2001US-029959P.
23-JUN-2001, 2001US-039999P.
24-JUN-2001, 2001US-039999P.
25-JUN-2001, 2001US-03999P.
26-JUN-2001, 2001US-03999P.
27-JUN-2001, 2001US-03999P.
28-JUN-2001, 2001US-03999P.
28-JUN-2001, 2001US-0310S9P.
28-JUN-2001, 2001US-0310S9P.
28-JUN-2001, 2001US-0310S9P.
28-JUN-2001, 2001US-0310S9P.
28-JUN-2001, 2001US-0310SP.
28-JUN-2001, 2001US-03199P.
28-JUN-2001, 2001US-03199P.
28-JUN-2002, 2002US-0359964P.
28-JUN-2002, 2002US-0379444P.
             04-JUN-2002; 2002WO-US017443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-210149/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or CNS diseases.
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Kekuda R;

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The invention relates to novel isolated polypeptides, mature form of the polypeptide, a sequence that is 95% identical to the polypeptide or the polypeptide comprising one or more conservative substitutions. The NOVX polypeptide is useful for treating or preventing a pathology associated with the polypeptide e.g. disorders associated with aberrant expression or activity of the polypeptide, such as cancer, diabetes, obssity, and candocrine, CNS and inflammatory disorders. They can also be used in various detection and screening assays, chromosome mapping, tissue typing and predictive medicine. This sequence corresponds to a primer used to amplify and isolate the coding sequence for one of the polypeptides of

Example B; SEQ ID NO 535; 772pp; English.

Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;

the invention.

schultz621-3.rng

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Gaps

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Indels

4

Mismatches

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Conservative
                                                                                                                                                                                                   Homo sapiens.
Synthetic.
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                                                                                                                                                                                                                                                                                                                                                         01-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                 20-MAY-1999;
                                                                                                                                                                                                                                                                                                         27-FEB-2003.
16;
                                                                                                      AAD58980;
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(COOK/)
(TILL/)
(HARD/)
                                                                                                                                                                                                                                                                                                                                                                                                                        (ECKE/)
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 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inhibiting corneal allograft rejection, by contacting an allograft with a formulation having an oligonucleotide targeted to intercellular adhesion molecule-1, extracellular adhesion molecule-1 or vascular cell adhesion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of inhibiting corneal allograft rejection, by contacting the allograft with a topical formulation comprising an antisense oligonucleotide targeted to intercellular adhesion molecule-1 (ICAM-1), extracellular adhesion molecule-1 (ICAM-1), extracellular adhesion molecule-1 (WCAM-1). The oligonucleotide is useful for inhibiting corneal allograft rejection or for preserving a corneal explant ex vivo, where the explant is human. This sequence
                                                                                                                                                                                                                                                                                       /*tag= a
/note= "all internucleotide linkages are phosphodiester
bonds"
                                                                                                                                                                                         ss; primer; immunosuppressive; antisense therapy; corneal allograft rejection; intercellular adhesion molecule-1; ICAM-1; extracellular adhesion molecule-1; BLAM-1; vascular cell adhesion molecule-1; VCAM-1; corneal explant.
                                                                                                                                                                                                                                                                                                                   1. .20
/*tag= b
/mod base= CTHER
/note= "OTHER = all A, C and U are 2'-fluoro bases or o-methyl"
                             Gaps
                             ö
         0.8%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 9.3e+02;
iive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   corresponds to one of the oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; SEQ ID NO 15; 106pp; English.
                                                                                                                                                                                                                                                                     Location/Qualifiers
                                               TCAAGCIGGCIGACTITGGC 1041
                                                                                                                                                                        Human ICAM-1 targeted primer #15.
                                                            TGAAGATTGCTGACTTCGGC 1
                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                   18-OCT-2001; 2001US-00982262.
                                                                                                                                                                                                                                                                                                                                                                                                                 16-OCT-2002; 2002WO-US033236
                                                                                                                ADC38989 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mirabelli CK;
                                                                                                                                                    (first entry)
(ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-403142/38.
                                                                                                                                                                                                                                                                               misc difference
                                                                                                                                                                                                                                                                                                                                                                            WO2003032920-A2
                                                                                                                                                                                                                                                                                                                     modified base
                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bennett CF,
                                                                                                                                                     18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                             24-APR-2003
                                                                                                                                                                                                                                          Synthetic
                                                                                                                                   ADC38989;
                                                                20
                                                                                              RESULT 1418
                                                                                                       ADC38989,
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0.8%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 9.3e+02;

Best Local Similarity

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Formulation, useful for treating inflammatory bowel disorder, e.g. ulcerative colitis or Crohn's disease, comprises oligonucleotide for rectal delivery.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ë
                                                                                                                                                                                                                                                                                                                                                                                              Inflammatory bowel disorder; ulcerative colitis; Crohn's disease; cellular proliferation; intracellular adhesion molecule; ICAM-1; phosphorothioate backbone; antisense; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Manoharan
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80.0%; Pred. No. 9.3e+02;
ative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hardee GE,
                                                                                                                                                                                                                                                                                                                                                  Human ICAM-1 antisense oligo, ISIS 1939.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cook PD, Tillman L,
226 GAGAGTGGTGGTGGTGGCGG
                                                 20 chcheceenacreereece
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/mod_base=
                                                                                                                                                                                     AAD58980 standard; DNA; 20
                                                                                                                                                                                                                                                                                                 18-DEC-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TENG C.
COOK P D.
TILLMAN L.
HARDEE G E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-596370/56.
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MANOHARAN M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified base
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AAD59446;

RESULT 1420

a ò

AAD59446

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New antibody that binds to an epitope of an inducible human phosphofructokinase-2 isozyme, useful for diagnosing or treating cancer, inflammation or cachexia.
                                                 Cytostatic; immunomodulator; phosphofructokinase isozyme; iPFK; cancer; inflammation; cachexia; enzyme linked immunosorbant assay; ELISA; therapy; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to an isolated antibody that binds to an epitope of an inducible human phosphofructokinase-2 (iPFK-2) isozyme. T antibody is useful for treating cancer, inflammation and cachexia. The antibody can also be used in enzyme linked immunosorbant assay (ELISA) immunological assays. The present sequence is S-iPFK-2 sense phosphorothioate oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA vaccine; flatfish rhabdovirus; HIRRV; fish; immunity;
transcriptional-control; cytomegalovirus immediate-type promoter;
immunogenic; virucide; gene gun; ss; primer.
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0.8%; Score 13.6; DB 1; Length 20;
Best Local Similarity 80.0%; Pred. No. 9.3e+02;
Matches 16; Conservative 0; Mismatches 4; Indels 0
                 S-iPFK-2 (A) sense phosphorothicate oligonucleotide.
                                                                                                                                                                                   1. .20 a
/tag= a
/mod_base= OTHER
/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mitchell RA;
                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                (PICO-) PICOWER INST MEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Col 10; 31pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                           97US-00961578.
98US-00183846.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chesney JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-743054/70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hirame rhabdovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JP2003155254-A.
                                                                                                                                                                   Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                         31-OCT-1997;
30-OCT-1998;
                                                                                                                              Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD22540;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antibody that binds to an epitope of an inducible human phosphofructokinase-2 isozyme, useful for diagnosing or treating cancer,
                                                                                                                                                                                                                                                                                  cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to an isolated antibody that binds to an epitope of an inducible human phosphofructokinase-2 (IPFK-2) isozyme. T antibody is useful for treating cancer, inflammation and cachexia. The antibody can also be used in enzyme linked immunosorbant assay (ELISA) immunological assays. The present sequence is AS-iPFK-2 antisense phosphorothicate oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                      Cytostatic; immunomodulator; phosphofructokinase isozyme; iPFK; cainflammation; cachexia; enzyme linked immunosorbant assay; ELISA; therapy; phosphorothioate; antisense; ss.
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                                                                                                                                                                                                                                          AS-iPFK-2 (A) antisense phosphorothioate oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "Phosphorothioate backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chesney JA, Mitchell RA;
                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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226 GAGAGTGGTGGTGGCGG 245
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/mod_base= OTHER
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                   20 GAGAGGGAAGTGGTGGGGG 1
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                                                                                                                              BP
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98US-00183846.
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AAD59445 standard; DNA; 20
                                                                                                                              AAD59446 standard; DNA; 20
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                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inflammation or cachexia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-743054/70.
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modified_base
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                                                                                                                                                                                                     18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                     Unidentified
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AAD59445;

BXXXE

RESULT 1421 AAD59445/c

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Best Loc Matches

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Gaps . 0

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A typing method for single nucleotide polymorphism (SNP) of several hundred thousands of SNP sites with comparatively a small amount of
                                                                                                                                                                                                                                           (RIKA ) RIKAGAKU KENKYUSHO
                                                                                                                                                                                                                                                   WPI; 2003-397221/38.
                                                                                                                                                                                                               JP2002300894-A.
                                                                                                                                                                                 15-JAN-2004
                                                                                                                                                                                                                      LS-OCT-2002.
                                                                                                                                                                                                                                                                 genome DNA.
                                                                                                                                                                          ADD68463;
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                                                                                                                                                            RESULT 14
ADD68463
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This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (Lohl) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (Asi) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somaticels of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene present in a population and targets only one allelic forms present in the normal somatic cells, and the first gene. The products and methods can be
                                           The invention relates to a novel method for typing a single nucleotide bolymorphism (SNP) using a small amount of genomic DNA comprishing simultaneous amplification of plural base sequences containing one or more SNP sites and differentiation of the bases within the SNP sites. The method of the invention may be useful for typing several hundred thousand SNP sites using only a comparatively small amount of genomic DNA. The current sequence is that of the SNP typing-related PCR primer of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premaligant metaplastic lesion; endometriosis; dyphastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                        Score 13.6; DB 1; Length 20;
Pred. No. 9.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                             Sequence 20 BP; 6 A; 6 C; 6 G; 2 T; 0 U; 0 Other;
Example 2; SEQ ID NO 20; 45pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 7; 605pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                765 GCTCAAGGACCTCAAACACG 784
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human polymorphic region 291.
                                                                                                                                                                                                                                                                                                                              0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Housman D, Ledley FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-521232/44
                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                               invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ26102;
                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel DNA vaccine for flatfish rhabdovirus (HIRRV) infected fishes, which provides immunity against HIRRV. The vaccination method uses a DNA construct comprising a transcriptional-control sequence containing cytomegalovirus immediate-type promoter, operably coupled to a nucleotide sequence encoding an immunogenic polypeptic of HIRRV. The DNA vaccine has virucide activity. The HIRRV DNA vaccine is useful for administering to a fish belonging to the flatfish family by gene gun. The HIRRY DNA vaccine is useful for inducing preventing HIRRV infection in flatfish. The HIRRY DNA vaccine is effective in enhancing immunity of fish infected by HIRRV. This polymucleotide sequence represente an oligo used in the analysis of the mRNA expression level from the muscles of flatfish, following an innoculation with the flatfish rhabdovirus vaccine of the invention.
                                                                                                                                                                                                                                                                       DNA vaccine for flatfish rhabdovirus infected fishes has DNA construct comprising a transcriptional control sequence coupled to a nucleotide sequence encoding an immunogenic protein of flatfish rhabdovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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80.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                            Example 6; Fig 5; 13pp; Japanese.
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                                                                                                                                             (MEIJ ) MEIJI SEIKA KAISHA LTD (AOKI/) AOKI H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     B
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                     26-SEP-2001; 2001JP-00294473.
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                                                                    06-SEP-2001; 2001JP-00271068.
10-SEP-2001; 2001JP-00274202.
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Best Local Similarity 80.0'
---nes 16; Conservative
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Nucleic acid sequence single primer amplification - useful for genomic variation analysis and polymorphism detection for restriction fragment length data.
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correlations to diseases. The present sequence is an example of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match

0.8%; Score 13.4; DB 1; Length 15;
Best Local Similarity 93.3%; Pred. No. 7.7e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                     0.8%; Score 13.6; DB 1; Length 21;
80.0%; Pred. No. 9.7e+02;
7ative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP; 5 A; 10 C; 0 G; 0 T; 0 U; 0 Other;
                                                              Sequence 21 BP; 3 A; 6 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Single primer amplification; SPAR; ss.
                                                                                                                                                                                               1459 TTCCTCAGTCTGGGGGAGCG 1478
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(first entry)
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(first entry)
                                                                                                     Query Match
Best Local Similarity 80.0 Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic primer (261)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1992-183683/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9207948-A1
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19-NOV-1992
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18-APR-1997
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                                                                                                                                                                                                                                                                                                                                                                                               AAQ24934;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype
                    cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic lesions, benign tumours, endometricsis, polycystic kidney disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular diesase; coronary artery disease; forenaics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
    used in the diagnosis, prevention and treatment of LOH disorders, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mccarthy JJ;
                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /standard_name= "single_nucleotide_polymorphism"
                                                                                                                                                                                                                                           ö
                                                                                                                                                                                             0.8%; Score 13.6; DB 1; Length 21;
80.0%; Pred. No. 9.7e+02;
tive 0; Mismatches 4; Indels
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                                                                                                                                                   Sequence 21 BP; 1 A; 5 C; 10 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human gene single nucleotide polymorphism #2298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ireland JS, Bolk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
replace(11,G)
                                                                                                                                                                                                                                                                                       1659 CACCCCTCACAGGGCAGCCC 1678
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                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF97537 standard; DNA; 21
                                                                                                                                                                                                                      Best Local Similarity 80.0
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-226749/23.
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Variation

AAF97537;

RESULT 1425 AAF97537

Query Match

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888888888

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Gaps

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Enzymatic nucleic acid, ribozyme, trans cleavage; inhibition; gene expression, downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; INF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease; atheroselerosis; myocardial infarction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS;
Human relA hammerhead ribozyme target sequence (nt. position 631).
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Homo sapiens

WO9523225-A2

31-AUG-1995

95WO-IB000156 94US-00201109 23-FEB-1995;

94US-00291433. 94US-00292620. 94US-0029320. 94US-00300000. 94US-00311486. 94US-0031149. 94US-00314397. 94US-00224483. 94US-00227958. 94US-00228041. 94US-00319492. 94US-00321993. 94US-00334847. 94US-00337608. 94US-00245736 94US-00271280 94US-00345516 94US-00291932 07-OCT-1994; 11-OCT-1994; 04-NOV-1994; 08-SEP-1994; 28-NOV-1994; 28-SEP-1994; 18-MAY-1994 06-JUL-1994 23-SEP-1994 10-NOV-1994 02-SEP-1994

(RIBO-) RIBOZYME PHARM INC.

94US-00363233 95US-00380734 DT, Chowrira B, Direnzo A, Draper KG, Dudycz LW; Karpelsky A, Kisich K, Matulic-Adamic J, Mcswiggen JA; Pavco P, Belgleman I, Sullivan SM, Sweedler D, Thompson JD; Usman N, Wincott FE, Woolf T; Stinchcomb DT, Grimm S, Modak A, Tracz D,

WPI; 1995-351090/45

use Ribozymes having modified bases and methods for producing them - for in inhibiting disease related genes.

Claim 2; Page 228; 407pp; English.

The present sequence represents a preferred target sequence for an enzymatic nuclecided acid (i.e. a ribozyme) which cleaves relA mRNA at the nucleotide base position indicated in the DE line. The reAA gene product is a subunit of the transcriptional regulator NF-kappaB and is implicated specifically in the induction of inflammatory responses. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their

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                 sequences and thereby inhibit relA expression, making them potentially useful for treating rheumatoid arthritis, restenosis and asthma as well as for increasing tolerance to transplanted tissues. The potential immunosuppressive properties of a ribozyme that cleaves relA mRNA means that uses are limited to local delivery, acute indications or ex vivo treatment. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                    Gaps
  nuclease resistance. The ribozymes are designed to cleave the target
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                                                                                                                                                                                                                                  0.8%; Score 13.4; DB 1; Length 15; 66.7%; Pred. No. 7.7e+02; ive 4; Mismatches 1; Indels
                                                                                                                                                                                  Sequence 15 BP; 4 A; 5 C; 1 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                   Query Match
Best Local Similarity 66.75
Warches 10; Conservative
                                                                                                                                                                                                                                                                                                                               539 CCATCTTTGACAAGC
88888888888
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1 CCAUCUUUGACAAUC 15

AAX75669 standard; RNA; 15

AAX75669;

(first entry) 28-JUL-1999

Human flt-1 and KDR hammerhead ribozyme target site #3.

Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.

Homo sapiens

WO9715662-A2

01-MAY-1997.

96WO-US017480, 25-OCT-1996; 95US-0005974P. 26-OCT-1995; 11-JAN-1996;

(RIBO-) RIBOZYME PHARM INC. (CHIR) CHIRON CORP.

Stinchcomb D, Pavco P, Mcswiggen J,

Escopedo

WPI; 1997-259017/23

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.

Example 9; Page 191; 218pp; English.

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vaccular endothelial growth factor (WRGF). A patient (preferably human) having a condition associated with the level of the fms-like tyzosine kinase 1 (EL-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule arthritis) can be vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention

Sequence 15 BP; 7 A; 1 C; 3 G; 0 T; 4 U; 0 Other;

Query Match

DB 1; Length 15; 0.8%; Score 13.4;

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The present sequence represents a hybridisation probe used to differentiate between pathogenic and vaccine strains of cattle brucellosis. The method comprises digestion of DNA from the test strain with restriction enzyme Nco 1, transfer of the fragments obtained to filters, subsequent fixing of these onto the filters, hybridisation with a labelled sample, and examination of the results. (Updated on 25-WAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                           Differentiating pathogenic and vaccine strains of cattle brucellosis -
using restriction digestion with Nco 1 and transfer of the DNA fragments
to filters in an electric field.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tag sequence; colorectal cancer; pancreatic cancer; colon cancer; diagnosis; prognosis; treatment; ss.
              Hybridisation probe; differentiation; pathogenic; vaccine strain; cattle brucellosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  / Match 0.8%; Score 13.4; DB 1; Length 15; Local Similarity 93.3%; Pred. No. 7.7e+02; nes 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 5 A; 10 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                           Mullakaev OT;
                                                                                                                                                                                                                                        (KZVE=) KAZAN VETERINARY MED ACAD.
                                                                                                                                                                                                         94RU-00024845.
                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Col 8; 4pp; Russian.
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                                                                                                                                                                                                                                                                           Idrisov GZ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-MAY-1999 (first entry)
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                                                                   Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Differentiating pathogenic and vaccine strains of cattle brucellosis -
using restriction digestion with Nco 1 and transfer of the DNA fragments
to filters in an electric field.
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                  Gapa
                                                                                                                                                                                                                                                                                                                        Hybridisation probe, differentiation, pathogenic, vaccine strain, cattle brucellosis; ss.
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93.3%; Pred. No. 7.7e+02;
tive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                       DNA sequence of the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (KZVE=) KAZAN VETERINARY MED ACAD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure, Col 4; 4pp; Russian.
                                                                                                                                                                      BP.
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                                                                                  15 ATTTCCATATTTGCA 1
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(first entry)
                                                                                                                                                                                                                                        (revised)
(first entry)
Best Local Similarity 93.3
Matches 14; Conservative
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16-OCT-1998
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16-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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Gaps

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Use of isolated gene transcripts - useful for developing products for diagnosis, prognosis and treatment of cancers, particularly colon and pancreatic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Extracting sequences of bases from dideoxyribonucleic acid templates research and medical applications, involves creating a new set of molecules which introduce error correcting code, from the template.
                                                                                                                                                                                                                                                                                                                                                                                                                   Dideoxyribonucleic acid; dDNA; research; medical application;
                                                                                                                                                                                                        Sequence 15 BP; 4 A; 3 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                               Original DNA template oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                              data communication; DNA sequencing; ss.
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                                               Claim 2; Page 34; 120pp; English.
                                                                                                                                                                                                                                                                   926 TCCAGCTGCTCCGTG 940
                                                                                                                                                                                                                                                                                                                                      AAA92356 standard; DNA; 15
                                                                                                                                                                                                                                                                                   TCCAGCTGCTCCATG 1
                                                                                                                                                                                                                                                 14; Conservative
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Best Local Similarity
Matches 14; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-587794/56.
                                                                                                                                                                                        treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (DAVI/) DAVIES
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(first entry)

98CA-02256128.

98CA-02256128

BP.

0.8%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 7.7e+02; tive 0; Mismatches 1; Indels

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weethod (I) has been developed for generating, in the same reaction vessel, a selected set of codons (II). The method comprises providing two (optionally three) sets of monouncleosides, mononucleotides.

(optionally three) sets of these and optionally repeatedly adding a third set, where (II) includes at least one codon having A or G at the third set, where (II) includes at least one codon having A or G at the third set, where (II) includes at least one codon having A or G at the third set, where (II) includes at least one codon having a or a stop codon. Also described is a method (II) for generating an oligomucleotide from (II), comprising the method (II), followed by repeating the method or thin an oligomucleotide of the desired length is achieved. (I) and (II) are useful in care useful in particular the methods are useful in pharmaceutical research. In particular the methods are useful in pharmaceutical research. In particular the methods are useful in techniques, ribosome display techniques and protein-mucleic acid fusion techniques. Codon-randomised DNA can also be used in cellular cultures (In vivo) for protein expression, or for in vitro applications using, e.g. T7 RNA polymerase, and in vitro translation systems. The present sequence represents an oligomucleocide which is used in the cemplification of the present invention
                                                                                                                                                                                                                                           ö
obtain accurate sequence estimate. (I) is useful for a research and medical applications. (I) minimises extor rates in sequencing or testing nucleic acids. The present sequence represents an original DNA template which is used in the exemplification of the present invention
                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Acid/base orthological deprotection scheme 15-mer oligonucleotide #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Acid/base orthological deprotection scheme; DNA synthesis; codon randomised nucleic acid; randomised cassette mutagenesis; phage display, ribosome display; protein-nucleic acid fusion; protein expression; in vitro translation system; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthesis of selected codon randomized nucleic acids useful for generation of DNA or RNA sequences for pharmaceutical research.
                                                                                                                                                                                                                                           .
0
                                                                                                                                                                                  0.8%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 7.7e+02; vative 0; Mismatches 1; Indels
                                                                                                                                  Sequence 15 BP, 5 A, 4 C, 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 3 A; 5 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 8; Page 29; 61pp; English.
                                                                                                                                                                                                                                                                                           1326 CAAGTACCGAGCCGA 1340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA29402 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-0102299P.
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                                                                                                                                                                                                                                                                                                                                              1 caagraccaagcrda 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-AUG-2000 (first entry)
                                                                                                                                                                 Query Match
Best Local Similarity 93.3%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lohse P, Kuimelis RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-293102/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHYL-) PHYLOS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200018778-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA29402;
                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1433
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA29402
8888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX30947-31815 represent tag sequences of transcripts that are differentially expressed in colorectal cancer, in pancreatic cancer, or in both. The tag sequences can be used to identify genes by matching the tag to a gen data base member, or by using the tag sequences as probes to isolate unidentified genes from cDNA libraries. The tag sequences can also be used in a method for diagnosing colon or pancreatic cancer in a sample suspected of being neoplastic. The method comprises comparing the level of at least one transcript in a first sample of a tissue to a second sample, where the first sample is a colonic tissue suspected of being neoplastic and the second sample is a normal human colonic tissue. The transcript is identified by a tag selected from AAX30947-31815, The methods of the invention can be used in the diagnosis, prognosis and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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1684 TACATCTTCCCTGCT 1698

15 racarrrcccrecr

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AAF46589 standard; DNA; 15

30-MAR-2001 (first entry) IGFBP3 oligonucleotide #9

AAF46589;

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Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; Insulin-like Growth Factor I receptor; IfFF1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hypermeovascular condition; hyperplamis; kelmosis; sclerotic disease; necovascular condition; hyperplamis; selence is clerotic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-19516). The method is useful for ameliotating the effects of psoriasis, ichthyosis, pityriasis, ruba, pitaris, serborrhoea, keloids, keratosis, chophanias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                          Gaps
                                          ..
  Length 15;
0.8%; Score 13.4; DB 1; Length 1
33.3%; Pred. No. 7.7e+02;
tve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Edmondson SR;
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                                                                                                                                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0140345P
                                                                              374 AGGCTTCAGCCACGT 388
                                                                                                                                                                                                                                                                                                                                         IGF-I oligonucleotide #1371.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUN-2000; 2000WO-AU000693
                                                                                                                                                                                                                    AAF50411 standard; DNA; 15
                                                                                                                                                                                                                                                                                                (first entry)
                   l Similarity 93.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wraight CJ, Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-041421/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-DEC-2000.
                                                                                                                                                                                                                                                        AAF50411;
                     Local
                   Best Loca
Matches
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Wraight CJ, Werther GA, Edmondson SR;

WPI; 2001-041421/05.

(MURD-) MURDOCH CHILDRENS RES

21-JUN-2000; 2000WO-AU000693.

WO200078341-A1 Homo sapiens.

99US-0140345P.

21-JUN-1999;

Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding proctein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilatis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keartosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

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Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Action of the present inversions. The method comparises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense oligonuclectides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serbornhoea, keloids, keratosis, inchthyosis, pityriasis, varts, benign growths, canners of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a method for ameliorating the effects of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15 BP; 1 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1634 GCAGGCAGCGGCTGG 1648
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammation.
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Gaps

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0.8%; Score 13.4; DB 1; Length 15; larity 93.3%; Pred. No. 7.7e+02; Conservative 0; Mismatches 1; Indels

Local Similarity les 14; Conserv

Query Match Best Local Si Matches 14;

Sequence 15 BP; 7 A; 1 C; 5 G; 2 T; 0 U; 0 Other;

vessels or any other hyperplasia

AAF50702 standard; DNA; 15 BP.

IGF-I oligonucleotide #1662

(first entry)

30-MAR-2001

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisomediacedride, (for Insulin-like Growth Factor [Gre]-1 creeptor, IGF binding protein [IGFB]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation.

Inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the affects of psoriasis, oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, neoplasias, soleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, the brain or skin, growth factor—mediated malignancies, other sclerotic disease, kidhey disease, hyperproliferation of the inside of blood vessels or any other hyperplasia
                                                                                                                                                                            Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF1; pityriasis; IGF binding procein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; necovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15 BP; 7 A; 1 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MURD-) MURDOCH CHILDRENS RES INST.
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               AAF50410 standard; DNA; 15 BP
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                                                                                                                                          IGF-I oligonucleotide #1370
                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inflammation.
                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                  30-MAR-2001
                                                          AAF50410;
AAF50410,
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Edmondson SR;

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                                Gaps
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  Length 15;
                            1; Indels
O.8%; Score 13.4; DB 1;
Similarity 93.3%; Pred. No. 7.7e+02;
14; Conservative 0; Mismatches 1;
                                                      1685 ACATCTTCCCTGCTT 1699
                            14; Conservative
Query Match
Best Local S
                            Matches
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RESULT 1437 AAF50702/c

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Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic, dermatological, cardiant, virucide, ophthalmological, keloid, skin disorder, insulin-like Growth Factor. I receptor; IGF-1, pitrylianis; IGF binding procein, IGFBP-2, IGFBP3, inflammation, psoriasis, pilaris; growth factor mediated cell proliferation, ichthyosis, serborrhoea; ruba, keratosis, neoplasia, scalaroderma, wart, skin cancer; sclerotic disease; hypermeovascular condition, hyperplasia, kidney disease; necovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wraight CJ, Werther GA, Edmondson SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MURD-) MURDOCH CHILDRENS RES INST.
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                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-DEC-2000.
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Ameliorating the effects of a disorder, e.g. psoriasis, by administering V (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

The present invention relates to a method for ameliorating the effects of antisense oligonucleotide, (for Insulin-like Growth Factor [IGF] receptor, IGF binding protein [IGFBP] - or IGFBP], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present esquence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153 oligonucleotides of the present invention (see AAF45151 and AAF45153 oligonucleotides of the present invention (see AAF45151 and AAF45153 olifothyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, ineoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic meneral or strain and present of the inside of blood Example 8; Page 71; 201pp; English. vessels or any other hyperplasia

ö Gaps ô 0.8%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 7.7e+02; ative 0; Mismatches 1; Indels 1283 CAGGCATCCTGTCCA 1297 Local Similarity 93.3 es 14; Conservative Query Match Matches ઠે

Sequence 15 BP; 2 A; 3 C; 7 G; 3 T; 0 U; 0 Other;

ABZ34171 standard; DNA; 15 BP. 15 caggcarccrecca RESULT 1438 ABZ3417 셤 RXXX

ABZ34171;

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The present invention describes a method for detecting mutations associated with anti-HV drug resistance in a patient by detecting at least one of the mutations K103N/R, V106A/IL, Y181CI, M184V/I, Y181E, C190A/S/R, T215Y/F/D/S/A and/or O151M/L in the reverse transcriptase (RT) of HV strains in a biological sample using a specific set of probes optimised to function together in a reverse-hybridisation assay. The method and the nucleic acid sequences used in the method are useful for determining viral mutations and/or polymorphisms in the HIV RT gene associated with resistance. The probes are useful for the genetic detection, preferably in vitro detection of the mutations K103N/R, V105M/IL, M181C/I, V181C/I, O151M/L, M184V/I, Y181E, G190A/S/R and/or C151SY/F/D/S/A in the RT of HIV strains in a biological sample, where the mutation is associated with anti-HIV drug resistance. The method provides a rapid, reliable and precise assay or determination and monitoring of antiviral drug resistance or mutations associated with drug resistance of viruses containing RT genes. ABZ34642 represent HIV RT contains and probes which are used in the exemplification of the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting mutations associated with anti-HIV drug resistance comprises detecting at least one of the mutations in the HIV reverse transcriptuse gene by using probes optimized to function together in a reverse-hybridization assay.
                                                                   Human immunodeficiency virus; HIV; reverse transcriptase; RT; enzyme; detection; mutation; anti-HIV drug resistance; polymorphism; resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gabe
                                    HIV-1 reverse transcriptase mutation detection probe SEQ ID NO:413.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 7.7e+02; tive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 4 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; Page 27; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human colon cancer SAGE tag #233
                                                                                                                                               Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                          11-JAN-2001; 2001EP-00870005.
20-APR-2001; 2001EP-00870085.
24-APR-2001; 2001US-0286102P.
                                                                                                                                                                                                                                                                         09-JAN-2002; 2002WO-EP000153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  867 GCAGTACCTGGATGA 881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 GCAGTACGTGGATGA 15
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ID ABK32132 standard; DNA; 15
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 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                    (INNO-) INNOGENETICS NV.
                                                                                                                                                                                                                                                                                                                                                                                                                        De Smet K, Stuyver L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-590680/63.
                                                                                                                                                                                                   WO200255741-A2
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   31-JAN-2003
                                                                                                                                                                                                                                      18-JUL-2002
                                                                                                            probe; ss.
                                                                                                                                                                  Synthetic.
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The invention relates to an isolated, purified human nucleic acid (I) that has the same sequence as a mRNA found in humans and is a SAGE (scrial analysis of gene expression) tag comprising a single stranded probe containing at least 10 consecutive nucleotides. SAGE tags, are diagnostic and prognostic markers of cancer, especially of the colon and pancreas. ABKX1900-ABKX2770 represent human colon and pancreatic cancer SAGE tags of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
            Human; colon cancer; colorectal cancer; pancreatic cancer; SAGE tag; estal analysis of gene expression; diagnostic; prognostic; probe; cancer marker; ss.
                                                                                                                                                                                                                                                                           as
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    16
    /*tag= a
    /note= "Linked via phosphorothioate linkages"

                                                                                                                                                                                                                                                                        New human nucleic acid containing specific SAGE tags, useful diagnostic markers for cancer, also derived probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rev response element; HIV isolate sf2; hybritope probe pool; hybritope mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 7.7e+02; ative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ineffective anti-HIV Rev response element probe 7819.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP; 4 A; 3 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                       Zhou W;
                                                                                                                                                                                                                       Zhang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                             Disclosure; Col 29; 161pp; English.
                                                                                                                                           98US-000B1646.
                                                                                                                                                                     98US-000B1646
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           926 TCCAGCTGCTCGTG 940
                                                                                                                                                                                                                       Vogelstein B, Kinzler KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT32677 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 rccagcrdcrccarg 1
                                                                                                                                                                                             SNINGO NINU ( OLYU)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 93.3*
Matches 14, Conservative
                                                                                                                                                                                                                                               WPI; 2002-153821/20.
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modified_base
                                                                                                                                           20-MAY-1998;
                                                                                                                                                                     20-MAY-1998;
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                                                               Homo sapiens
                                                                                       US6333152-B1
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(ISIS-) ISIS PHARM INC
(CHIR ) CHIRON CORP
             WPI; 1996-287198/29.
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modified_base
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13-MAR-1996
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                                                                                                                                                                                                           15-AUG-1995
      Collins ML;
                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                      3aker B,
                                                                                                                                            AAT11976;
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                                                                                                                                  AAT11976,
XBX1XBXBXBXBX1111X8XXBX1116XBX
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                                                                                                                              AAT11971-84 are antisense oligonucleotides (ONS) against human expressed vertivities of at least 50 % of control (ISIS 2922 shown in APT11961). It was found that up to 4 internal mismatches could be tolerated without loss of antiviral activity. Antisense ONS targeting CMV DNA or RNA coding for the ISI, ISE or DNA polymerase proteins have been shown to be effective in therapy, prophylaxis and diagnosis of CMV infection. The ONS may be modified to prophylaxis essistence and to increase their efficacy. Modifications include phosphorothicate backbones, alkyl and halogen-substituted sugar modelies at the 2' position. (Updated on 25-MAR-2003 to correct PF
New oligo-nucleotide inhibits cytomegalovirus replication - by binding to a portion of cytomegalovirus RNA, for the diagnosis, prophylaxis and treatment of CMV diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New peptide nucleic acid oligomers hybridisable to cytomegalovirus or papilloma:virus - are stable anti:sense molecules with high affinity for single stranded DNA, used for treating infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New oligomers are claimed which (A) have at least one peptide nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide nucleic acid, PNA, cytomegalovirus, CMV, papillomavirus; antiviral; diagnostic, 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; tive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide nucleic acid targetting CMV IE2 nuc sig 2.
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 0 A; 5 C; 3 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mirabelli CK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                       Example 10; Col 17; 66pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      135 GAAGAAGATCAAACG 149
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
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Best Local Similarity
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misc_feature
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AAT01678/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                              The sequences given in AAT32673-76 represent effective, and those in AAT32673-76 represent effective, and those in AAT32673-783 ineffective, anti-HIV Rev response element probes isolated from a hybritope probe pool. Hybritope mapping describes a method of determining superior sites for binding oligonucleotides to a target sequence, to identify improved discontinuous probes with high binding constants. The method comprises obtaining a series of oligonucleotides which are complementary to a known target sequence and which overlap each other by 1-4 mucleotides. Each of these sequences is contacted with the target sequence to permit specific hybridisation to determine presence or absence of specific hybridisation to determine oligonucleotides which bind within the known target sequence. This sequence was isolated using the probe sequences givne in AAT32670-72. The number of this probe corresponds to the 5' position on the HIV sf2 target to which the 3' end of the probe binds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense; cytomegalovirus; CMV; human; therapy; prophylaxis; diagnosis; intermediate early complex; IE1; IE2; DNA polymerase gene; ss.
                                                                                                                           Detecting target binding oligo-nucleotide(s) - using oligo-nucleotide probes with a nucleotide sequence which binds within a known sequence of a target nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 13.4; DB 1; Length 16; Best Local Similarity 93.3%; Pred. No. 8.2e+02; Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16 BP; 4 A; 5 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CMV antisense oligonucleotide (ISIS 5480).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                          Example 5; Page 27; 43pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT11976 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           156 GTCAATGACACTCCG 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
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acid (PNA) subunit and (B) have a sequence hybridisable to AUG region, 5 untranslated region, intron/exon (I/E) junction or coding sequence of curtanslated region, intron/exon (I/E) junction or coding sequence of hybridisable to the E, E2, E4, E5, E6, E7, L1 or L2 reading frames of a papillomavitus. The PNAs can be used to target RNA and single stranded DNA (scDNA) to produce antisense-type gene regulation moieties. Hence they may be used therapeutically for modulating cytomegalovirus and papillomavitus processes and also as diagnostics (e.g., as probes for specific maNAs). PNA oligomers have high affinity for complementary single stranded DNA. They are also able to form triple helices in which single strand binds with RNA or sabNA and a second RNA strand binds with the resulting double helix or with the first PNA strand binds with the resulting double helix or with the first PNA strand binds with the resulting double helix or with the first PNA strand change and are water soluble, which facilitates cellular upsake. Buther, shore they contain andes of non-blological amino acids, they are biostable and resistant to enzymatic degradation by processes. The present sequence targets CMV IE2 nuclear localisation

Sequence 17 BP; 0 A; 5 C; 3 G; 9 T; 0 U; 0 Other;

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Gapa
                             0
0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; Live 0; Mismatches 1; Indels
                                                       135 GAAGAAGATCAAACG 149
                                                                                  16 GAAGAGAGCAAACG 2
                              14; Conservative
                 Similarity
 Query Match
                 Local
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Human flt1 VEGF receptor hammerhead ribozyme substrate #474. BB AAX69179 standard; RNA; 17 (first entry) 28-JUL-1999 AAX69179; RESULT 1443 AAX69179/c

Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR, hammerhead ribozyme; hairpin ribozyme; detavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase i; kinase insert domain containing receptor; foetal liver kinase 1; ss

Homo sapiens

WO9715662-A2

96WO-US017480. 25-OCT-1996; 01-MAY-1997.

95US-0005974P. 26-OCT-1995; 11-JAN-1996;

(RIBO-) RIBOZYME PHARM INC. (CHIR) CHIRON CORP.

Escobedo J; Stinchcomb D, Pavco P, Mcswiggen J,

WPI; 1997-259017/23.

stability - useful for treating VEGF receptor(s) gene expression or mRNA rheumatoid arthritis, etc., in a human patient.

Claim 4; Page 61; 218pp; English.

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a MRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the

Length 17;

0.8%; Score 13.4; DB 1; 60.0%; Pred. No. 8.7e+02;

Query Match Best Local Similarity

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fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour anglogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; VBGF; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid archritis; coular disease; fms-like; tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid molecule modulating VEGF receptor(s) gene expression or
stability - useful for treating e.g. tumour angiogenesis, psoriasis,
rheumatoid arthritis, etc., in a human patient.
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0
                                                                                                                                                                   0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; rative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human KDR VEGF receptor hammerhead ribozyme substrate #483.
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                                                                                                                                  Sequence 17 BP; 7 A; 3 C; 3 G; 0 T; 4 U; 0 Other;
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96US-00584040.
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                                                                                                                                                                                                                                                                               16 Arriccararrigca 2
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                                                                                                                                                   Pavco P, Mcswiggen J,
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11-JAN-1996;
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Anti-CMV oligonucleotide #5480.
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AAX17893 standard; DNA; 17
                 (first entry)
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Best Local Similarity 93.35
Marches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                           Black IA, Woodbury D,
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-594570/50
                                                                                                                                    human; probe; ss
                                                                                                                                                                Synthetic.
Homo sapiens.
                 05-FEB-1999
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AAX17893/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes enzymatic nucleic acid molecules (NAMs) which specifically cleave RNA derived from an epidermal growth factor receptor (EGF-R) gene. AAV9721 to AAV98043 and AAV9899 to AAV9909090 represent specifically claimed target sequence from human EGF-R. AAV98044 to AAV98066 and AAV98867 to V9978 represent hammerhead ribozymes and hairpin ribozymes respectively for human EGF-R. The NAMs are useful for cleaving EGF-R RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit cell proliferation in the prevention or treatment of cancers. The NAMs can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enzymatic nucleic acids - which cleave RNA derived from an epidermal growth factor receptor, useful for inhibiting cell proliferation and for
                                                                                                                                                                                                                                               Human, epidermal growth factor receptor; EGFR; EGF-R; target sequence;
hammerhead ribozyme; hairpin ribozyme; inhibition; cell proliferation;
cancer; genetic drift; detection; mutation; ss.
   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.4; DB 1; Length 17;
80.0%; Pred. No. 8.7e+02;
tive 2; Mismatches 1; Indels
   Indels
                                                                                                                                                                                                                    Human EGF-R target sequence nucleotide position 2624.
   ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 8 C; 2 G; 0 T; 4 U; 0 Other;
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Akhtar S, Fell P, Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 5; Page 74; 109pp; English.
5
                                                                                                                               AAV97521 standard, RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV69694 standard; DNA; 17 BP.
                            1032 TGACTTTGGCCTGGC 1046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    989 CCCAGAACCTGCTCA 1003
                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US000730.
                                                                                                                                                                                                                                                                                                                                                                                                                               97US-0036476P
                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-00985162
                                                        3 véacurudécurides 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC. (UYAS-) UNIV ASTON.
                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 80.0
les 12; Conservative
9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-437449/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treating cancers.
                                                                                                                                                                                                                                                                                                                                          WO9833893-A2.
                                                                                                                                                                                                                                                                                                              sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                             31-JAN-1997;
04-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                 14-JAN-1998;
                                                                                                                                                                                         17-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                     06-AUG-1998,
                                                                                                                                                             AAV97521;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV69694;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1446
AAV69694
                                                                                                   RESULT 1445
                                                                                                                                                                                                                                                                                                              Ното
Matches
                                                                                                                    AAV9752
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Sequences AAV69693 and AAV69694 represent nested oligonucleotide probes corresponding to the exon 1 of the human glial cell line-derived neutrophic factor (GDNF) gene. These were used to identify the initiation of transcription of hGDNF gene. The invention relates to the use of the human GDNF promoter which contains a proximal section which cansures consistent low level GDNF expression in multiple cell types, and a distal section designed to alter transcription during development and in response to environmental stimuli. The GDNF promoter can be used for expressing GDNF in a cell, for identifying modulators and binding extrems of a GDNF promoter and modulators of GDNF expression. The products can be used for diagnosis and treatment of disorders involving GDNF such as neural degeneration, e.g. seizures, Parkinson's disease, Lou Gebrig GDNF during the prematal and neonatal stage. The GDNF promoter is also used for gene therapy and for expressing heterologous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated glial cell line-derived neurotrophic factor promoter - used to develop products for treating e.g. neuronal degeneration, immunodeficiency, haemophilia or proliferative disorders such as cancers.
                                                                                  GDNF, glial cell line-derived neurotrophic factor; promoter; seizure; transcription; environmental stimulus; modulator; neural degeneration; Parkinson's disease; Lou Gehrid's disease; developmental defect; tumour; gene therapy; neural degeneration; immunodeficiency; haemophilia; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ..
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Human GDNF gene exon 1 specific nested probe exon 1B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ramakrischnan L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genes for treating e.g. severe combined immunodefic proliferative disorders such as tumours and cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 17 BP; 2 A; 7 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schaar DG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 43; 69pp; English.
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New antisense oligonucleotides that target cytomegalovirus nucleic acid -
particularly including 2-methoxyethoxy sugar modifications, especially
for treating viral retinitis, with long-lasting retention in the retina.
                                                                                                                                                                                                                                                                                                                                                               Antisense oligonucleotides (AAX17861-X17924) are targeted to a nucleic acid (AAX1792-X17948) encoding IE (immediate early) 1 or 2, or DNA polymerase of cytomegalovirus (CWV) and are able to inhibit CWV replication. Optionally the oligonucleotides include at least one 2'-(2-methoxyethoxy) sugar modification or phosphorothioate internucleotide Inkages. The oligonucleotides are used to inhibit CWV infections (by in vivo or in vitro contact with cells, tissues or body fluids), especially to treat or prevent CMV infections, particularly retinitis
Antisense; oligonucleotide; immediate early; DNA polymerase; CMV; cytomegalovirus; inhibition; replication; sugar modification; phosphorothioate; infection; retinitis; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 0 A; 5 C; 3 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                Anderson KP, Chapman S,
                                                                                                                                                                                                                                                                                                                                         Claim 7; Page 30; 99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA21066 standard; RNA; 17 BP.
                                                                                                                                                  98WO-US006895.
                                                                                                                                                                            97US-00838715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUN-2000 (first entry)
                                                                                                                                                                                                                                Draper KG, Kisner DL,
                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                          WPI; 1998-568330/48.
                                                                   Human herpesvirus 5.
                                                                                                                                               07-APR-1998;
                                                                                             WO9845314-A1
                                                                                                                                                                            09-APR-1997;
                                                                                                                       15-OCT-1998
                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA21066;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Gaps
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Match 0.8%; Score 13.4; DB 1; Length 17; Local Similarity 93.3%; Pred. No. 8.7e+02; les 14; Conservative 0; Mismatches 1; Indels
                                                                                                   135 GAAGAAGATCAAACG 149
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16 GAAGAGAGCAAACG 2

Integrin alpha 6 subunit substrate sequence SEQ ID NO:4292.

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; amidiabetic factor; cytocatatic; antidabetic; ophthalmologic; antifainatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; veruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; ss.

Homo sapiens

W09950403-A2

07-0CT-1999

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The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl nuclear transporter (ARVT) gene, an integrin submit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene, AAA16775 to AAA17651 to AAA17651 to AAA17652 represent ribozyme sequences for ARNT, and AAA17651 to AAA17652 represent ribozyme sequences for AAA19087 to AAA11675 to AAA19185 to AAA19185 to AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18087 to AAA19155 to AAA191822 represent their corresponding target sequences; AAA19155 to AAA191822 represent their corresponding target sequences; AAA19155 to AAA21501 and AAA21501 to AAA21595 to AAA21500 and AAA21595 to AAA21501 and AAA21501 to AAA21595 to AAA21500 and AAA21505 to AAA21505 and AAA2150 and AAA22475 and AAA22475 to AAA23262, AAA23342 represent their corresponding target sequences; for integrin subunit beta 3, and AAA23476 to AAA23262, AAA23343 to AAA23472 represent their corresponding target sequences; the invention are used for modulating target sequences. The ribozyme of the invention are used for modulating target sequences. The ribozyme sepecially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as necular degeneration (ARMD), inflammation, and arthritis, as well as necular degeneration (ARMD), inflammation, and arthritis, as well as necular degeneration (ARMD), inflammation, and arthritis, as well as necular degeneration (ARMD), inflammation, and arthritis, as well as necular degeneration (ARMD), inflammation, and arthritis, as well as necular degeneration (ARMD), inflammation, and arthritis, as well as anglofibrome of tuberous scilerosis, pot-wine stains, sturge Weber syndrome, and other syndromes and diseases related to the where any and arthritis, and arthritis, and arthritis, and any and arthritis, and a
                                                                                                                                                                                                                                                                                                    Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors.
                                                                                                                                                                                  Coeshott C, Mcswiggen JA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               integrin subunit alpha-6, or integrin subunit beta-3
                                                                                                                                                                                                                                                                                                                                                                                              Claim 55; Page 185; 305pp; English.
                                                                                                                                                                                  Jarvis T,
   99WO-US006507.
                                                       98US-0079678P.
                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                               Pavco PA, Roberts E,
                                                                                                                                                                                                                                         WPI; 1999-591315/50.
24-MAR-1999;
                                                          27-MAR-1998;
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0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; ative 0; Mismatches 1; Indels Sequence 17 BP; 2 A; 0 C; 7 G; 0 T; 8 U; 0 Other; Local Similarity 93.3 Query Match Best Loca Matches

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Gaps 0

> 1314 ATACAACTACCCCAA 1328 16 ACACAACTACCCCAA

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RESULT 1449

257/c AAA23257 standard; RNA; 17 BP AAA23257;

19-JUN-2000 (first entry)

Integrin subunit beta 3 substrate sequence SEQ ID NO:6483.

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cyrostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot whne stain; Sturge Weber syndrome; ss.

Homo sapiens

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The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin alpha 6 subunit gene, and AAA1768 to AAA1761 to AAA17684 represent their corresponding target sequences; AAA17682 to AAA18885 and AAA19086 and AAA19152 to AAA1922 represent their corresponding target sequences; AAA17685 to AAA1982 to AAA1985 to AAA1986 and AAA19086 and AAA1985 to AAA1922 to AAA2160 and AAA19159 to AAA2169 and AAA19159 to AAA2169 and AAA2169 to AAA2169
                                                                                                                                                                                                                                                                                                                                                                                                            Novel ribozymes for modulating the synthesis, expression and/or stability
                                                                                                                                                                                                                                                                                                            Coeshott C, Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    integrin subunit alpha-6, or integrin subunit beta-3
                                                                                                                                                                                                                                                                                                                                                                                                                                            of an mRNA encoding an angiogenic factors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 54; Page 271; 305pp; English.
                                                                                                                                                                                                                                                                                                            Jarvis T,
                                                                                                                                         99WO-US006507.
                                                                                                                                                                                             98US-0079678P.
                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                         Roberts E,
                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-591315/50.
                                                                                                                                         24-MAR-1999;
                                WO9950403-A2
                                                                                                                                                                                          27-MAR-1998;
                                                                                 07-0CT-1999
                                                                                                                                                                                                                                                                                                      Pavco PA,
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Sequence 17 BP; 3 A; 4 C; 4 G; 0 T; 6 U; 0 Other;

Gaps ö / Match
0.8%; Score 13.4; DB 1; Length 17;
Local Similarity 93.3%; Pred. No. 8.7e+02;
les 14; Conservative 0; Mismatches 1; Indels Query Match Best Loca Matches

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RESULT 1450 AAA20471 AAA20471;

AAA20471 standard; RNA; 17 BP

19-JUN-2000 (first entry)

Integrin alpha 6 subunit substrate sequence SEQ ID NO:3697.

Human, aryl hydrocarbon nuclear transport, ARNT, TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme, angiogenic factor; cycostatic; antidabetic; ophthalmologic; antiinflammatory; antiathritic; antisoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;

The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl cleaving activity, which specifically cleave RNA encoded by an aryl of hydrocarbon nuclear transporter (ARNY) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA1760 and AAA1760 and AAA1760 to AAA1760 and AAA1760 and AAA1760 and AAA1760 and AAA1960 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA1900 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA1900 and AAA19155 to AAA19222 represent their corresponding target sequences; AAA19223 to AAA2180 to AAA2180 and AAA2180 to AAA2180 to AAA2180 and AAA2180 to AAA2180 to AAA2180 and AAA2180 to AAA2180 to AAA2180 to AAA2180 to CC AAA2180 to A Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors. Gaps tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; .. 0 Mcswiggen JA; 0.8%; Score 13.4; DB 1; Length 17; 33.3%; Pred. No. 8.7e+02; ve 6; Mismatches 1; Indels integrin subunit alpha-6, or integrin subunit beta-3 Sequence 17 BP; 2 A; 5 C; 3 G; 0 T; 7 U; 0 Other; Coeshott C, Claim 55; Page 147; 305pp; English. Jarvis T, 99WO-US006507 98US-0079678P 918 GITCCIGITCCAGCT 932 53.3%; (RIBO-) RIBOZYME PHARM INC. Local Similarity 53.3 Pavco PA, Roberts E, WPI; 1999-591315/50. WO9950403-A2 Homo sapiens 24-MAR-1999; 27-MAR-1998; 07-OCT-1999. Query Match Best Loca Matches ઠે 셤

AAA24802 standard; DNA; 17 BP. AAA24802; RESULT 1451 AAA24802

19-JUL-2000 (first entry)

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Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1300.

Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage;

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nucleic acids that interact, and optionally cleave, target sequences,
hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide;
gene expression modification; cancer; phosphorothioate; endonuclease;
anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                            Thompson JD, Beigelman L, Mcswiggen JA, Karpeisky A, Reynolds M, Zwick M, Jarvis T, Woolf T, Haeberli P,
                                                                                                                                                                                                                       Claim 77; Page 58; 148pp; English.
                                                                                     99WO-US008547.
                                                                                                             98US-00103636.
                                                                                                     98US-0082404P
                                                                                                                           (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                       used to treat cancer
                                                                                                                                                                             WPI; 2000-013248/01
                                                                                                                                                             Matulic-Adamic J;
                                  Homo sapiens
                                                 WO9954459-A2
                                                                                    19-APR-1999;
                                                                                                             23 - JUN - 1998;
                                                                                                    20-APR-1998;
                                                                  28-OCT-1999
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The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(dilthicate link, having endonuclease activity. (A), and more generally any catalytic link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A/) that modulates expression of the oestrogen receptor. C gene, are used to treat cancer (particularly of breast or endometrium). In vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of estrogen receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research resents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA2393 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences.

CC AAA24747 represent oestrogen receptor hammerhead ribozyme sequences.

CC AAA2593 to AAA26107 to AAA26218 represent their corresponding target sequences. and AAA26271 represent other ribozyme sequences and antennal and antisense oligonucleotides used in the exemplification of the present Sequence 17 BP; 5 A; 6 C; 4 G; 2 T; 0 U; 0 Other; Query Match Best Local Similarity

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Gapa
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0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; Live 0; Mismatches 1; Indels
                                    14; Conservative
                                  Matches
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1315 TACAACTACCCCAAG 1329 TACAACTACCCCGAG 16

AAF06373 standard; DNA; 17 BP AAF06373; RESULT 1452 AAF06373

Hammerhead ribozyme substrate #3170.

(first entry)

16-FEB-2001

Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.

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Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoletin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAAT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and
                                                                                                                                                                                                                                                                             Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 4 A; 5 C; 3 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                 Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                     Claim 42; Page 128; 164pp; English.
                                                                                                                                                                                                              Blatt L, Zwick M, Pavco P,
                                                                                                     11-APR-2000; 2000WO-US009721
                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                              WPI; 2000-647423/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                interferon alpha
                                 WO200061729-A2.
 Homo sapiens.
                                                                     19-OCT-2000.
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Bellon L;

Gaps .. Query Match 0.8%; Score 13.4; DB 1; Length 17; Best Local Similarity 73.3%; Pred. No. 8.7e+02; Matches 11; Conservative 3; Mismatches 1; Indels Query Match

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686 ACAACCTTGTGGCAC 700 ||| ||:|:||||| ACAUCCUUGUGGCAC 16 à ద

ABK03332 standard; RNA; 17 12-MAR-2002 (first entry) Human CD20 Inozyme #283. ABK03332;

RESULT 1453

ВЪ.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; noctropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; human immundeficioncy virus; HTV associated NHI; mantle-cell lymphoma; MCL; immunocytoma; IMC; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cenebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

WO200159103-A2. 16-AUG-2001. Synthetic.

Homo sapiens

09-FEB-2001; 2001WO-US004273

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 game and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids may be enzymatic nucleic acid cleaving a an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a CD20 in the presence of a divalent cation that is preferably Mg^2+, cofficient and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more the cation prophoma, leukaemia, B-cell lymphoma, low-grade or follicular non-cleakagin's lymphoma (NLH), bulky low-grade or follicular non-cleakagin's lymphoma (NLH), bulky low-grade or follicular non-cleakagin's lymphoma (NLH), bulky low-grade or follicular non-cleakagin's lymphoma (NLH), immunocytoma (INC), small B-cell lymphocytic lymphoma (NCL), immunocytoma (INC), small B-cell lymphocytic lymphoma (NCC) craseting nucleic acid may be contacted with a cell coleave (CC crase central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, and/or other neurodegenerative disease teates which respond to the modulation of NOGO expertation of NOGO expertation of NOGO expertation of NOGO disease, muscular disease, dementian in ninozyme of the invent
                                                                                                                                                                                                                                                                                                                                      Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence is an inozyme of the invention
                                                                                                                                                                                                                                                  Chowrira BM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 30; Page 150; 200pp; English.
                  11-FEB-2000; 2000US-0181797P.
28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                                                                                        RIBOZYME PHARM INC.
                                                                                                                                                                                                                                          Blatt L, Mcswiggen J,
                                                                                                                                                                                           CHOWRIRA B M.
                                                                                                                                            BLATT L.
MCSWIGGEN J.
                                                                                                                                                                                                                                                                                            WPI; 2001-607195/69
                                                                                                                                          (BLAT/)
(MCSW/)
(CHOW/)
                                                                                                                     (RIBO-)
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Sequence 17 BP; 4 A; 4 C; 6 G; 0 T; 3 U; 0 Other;

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Match 0.8%; Score 13.4; DB 1; Length 17; Local Similarity 93.3%; Pred. No. 8.7e+02; les 14; Conservative 0; Mismatches 1; Indels
  Query Match
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Matches
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ABK03331 standard; RNA; 17 BP ABK03331; RESULT 1454 ABK03331/ ID ABK0 XX AC ABK0 XX DT 12-M XX DE Huma

12-MAR-2002 (first entry)

Human CD20 Inozyme #282.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; intryme; Inozyme; Jeleaver; amberzyme; intryme; Inozyme; Jeleaver; amberzyme; Iymphoma; lumboma; lumboma; NHL; lymphoma; lekaemia; human immundeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

Homo sapiens. Synthetic.

WO200159103-A2.

16-AUG-2001.

09-FEB-2001; 2001WO-US004273

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

(RIBO-) RIBOZYME PHARM INC (BLAT/) BLATT L. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M. Chowrira BM; Blatt L, Mcswiggen J,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 30; Page 150; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a cD20 gene and a nucleic acid molecule which down control acids acids eaving a suboxyme or a regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids and NGH motif), a G-cleaver (cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) proposessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) proposessing an NCH motif). The CD20-targetting nucleic acid is used to cleave RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of cleaving RNA with a NGM with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of the cell and treat a patient having a condition associated with the level of CD20 in the teatment may further comprise the use of one or more the coll in particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Bodgkin's lymphoma (NRI), bulky low-grade or follicular NHI, lymphocytic lymphoma (NRI), timmunocytoma (IMC), small B-cell lymphocytic lymphoma, laukaemia, and inflammatory arthropathy. The NOGO targetting nucleic acid may be contacted with a cell to reduce NOGO gene in the presence of a divalent cation that is preferably Mg^2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO gene in the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more cell and treat a patient having a condition? Sociated with the level of the ROGOT The treatment may further comprise the use of one or more creates which respond to the modulation of sisease, ctaxia, Hunthigton's di

Human; mouse; otoferlin; OTOF; brain; auditory function; PCR primer; autosomal nonsyndromic prelingual deafness; DFNB9; ss.

Human otoferlin exon PCR primer #39.

(first entry)

13-FEB-2002

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Gaps

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AAS95074;

AAS95074 standard; DNA; 17

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The present sequence is a degenerate PCR primer used for amplifying the human cell cycle checkpoint protein, hohki DNA. The cell cycle checkpoint protein, hohki DNA. The cell cycle and timing of cell cycle transitions, and enaure that critical events such as DNA replication and chromosome segregation are completed with high fidelity. The chtl protein controls cell cycle in response to DNA damage. It functions as kinase and phosphorylates the key regulators of Cdk tyrosine phosphorylation. The checkpoint gene sequences are used as probes for a portion of the chromosome associated with tumours and other malignancies, as well as growth and/or development deficiencies. The chkl proteins are useful for generating specific antibodies and for inhibiting growth of
                                                                                                                                                                                                                                                                                                                                                                                    Human cell cycle checkpoint protein, hchkl DNA amplifying PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New Chkl proteins and gene sequences encoding the proteins useful as probes for a portion of the chromosome associated with tumors and other malignancies, growth and/or development deficiencies.
                                                                                                                                                                                                                                                                                                                                                                                                                         Human, cell cycle checkpoint; chkl; tumour; malignancy;
cell growth inhibitor; development deficiency; PCR primer; DNA damage;
kinase; ss.
                                                                       Query Match
0.8%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 8.7e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                      Sequence 17 BP; 5 A; 4 C; 6 G; 0 T; 2 U; 0 Other;
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     sequence is an inozyme of the invention
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                                                                                                                                                                                                                                                                           AAD03853 standard; DNA; 17 BP.
                                                                                                                                               396 TGAGGTGCAGTCTCC 410
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Novel human gene Otoferlin, underlying an autosomal recessive nonsyndromic prelingual deafness, DFNB9, and proteins encoded by the gene, implicated in deafness.

Claim 25; Page 17; 99pp; English.

Petit C;

El Amraoui A,

Cohen-Salmon M,

Yasunaga S, Grati M,

Weil D;

WPI; 2001-611499/70

(INSP) INST PASTEUR. (CNRS) CNRS CENT NAT RECH SCI.

23-MAR-2001; 2001WO-IB000578. 24-MAR-2000; 2000US-0191738P.

WO200170972-A2. Homo sapiens

27-SEP-2001.

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The invention relates to a purified polynuclectide (I) encoding a protein sequence (II) encoded by a novel human gene, otoferlin (OTOF) or the long human otoferlin isoform in brain. (I) was identified as underlying an autosomal nonsyndromic prelingual deafness DRNB9, and is thus useful for detecting deafness disease in humans and for characterising the functions of proteins and genes encoding them in auditory function. AAS95022-AAS95048 represent human and mouse otoferlin coding sequences, PCR
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muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 495 CCGCTGCCTGAGGG 509
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ABN08906 standard; DNA; 17
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0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; ive 0; Mismatches 1; Indels

1033 GACTTTGGCCTGGCC 1047

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Local Similarity es 14; Conserv

Best Loca Matches

Query Match

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Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:67.

(first entry)

29-MAY-2002

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 claim be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements and for case as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins or polymeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP-1 proteins, as specific biomolecule capture probes for surface-enhanced laser describing deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence captured in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polypeptide, for raising antibodies that recognize hdDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser descrption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234687P.
24-SEP-2000; 2000US-02356359P.
04-OCT-2000; 2000US-0236359P.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
                                                                               25-MAY-2001; 2001WO-US016981
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WO200192524-A2
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tch 0.8%; Score 13.4; DB 1; Length 17; al Similarity 93.3%; Pred. No. 8.7e+02; 14; Conservative 0; Mismatches 1; Indels
Query Match
Best Local S:
Matches 14
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165 ACTCCGAGGTGGCCG 179 ACTCGGAGGTGGCCG 1

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RESULT 1458 ABN00075/C ID ABN00075 standard; DNA; 17 BP. AC ABN00075;

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. 26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234667P.
27-SEP-2000; 2000US-0234667P.
04-OCT-2000; 2000US-0236339P.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667. 25-MAY-2001, 2001WO-US016981 (AEOM-) AEOMICA INC. WO200192524-A2 Homo sapiens 06-DEC-2001

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Shannon ME;

Hanzel DK, Rank DR, Chen W,

Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23. Disclosure, SEQ ID NO 67; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 claim be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids in samples, as amplification substrates to mucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for provide initial substrates for the recombinant engineering of hGDMLP-1 proteins or polypeptides may be protein, variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specific biomolecule and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The production and skeletal muscle disorders. hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1 may be used for chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this spatent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fig. The present invention.

Sequence 17 BP; 2 A; 5 C; 3 G; 7 T; 0 U; 0 Other;

Gaps .; 0 Query Match

0.8%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 8.7e+02;
Matches 14; Conservative 0; Mismatches 1; Indels

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New polypeptide, for raising antibodies that recognize hGDWLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDWLP-1.
                                                                                                                                                           Human, genome-derived myosin-like protein 1, GDMLP-1, hGDMLP-1, heart, muscle, myosin, chromosome 22, gene therapy, vaccine, heart disease, skeletal muscle disorder, amplicon, screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hanzel DK, Rank DR, Chen W, Shannon ME,
                                                                                                                                         Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:66.
                                                                                                                                                                                                                                                                                          26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000US-00264263.
30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
1181 ATGAGATGGCCACAG 1195
                                                                         ABN00074 standard; DNA; 17 BP
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                                                                                                                    (first entry)
            16 ATGAGATGGACACAG
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                                                                                                                    29-MAY-2002
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                                                                                              ABN00074;
                                                    RESULT 1459
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1
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The present sequence represents an oligomer used in the screening of the hGDWLP-1 sequence in the exemplification of the present invention, N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                      Query Match 0.8%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 8.7e+02; Matches 14; Conservative 0; Mismatches 1; Indels
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                                                                                                        Sequence 17 BP; 2 A; 5 C; 2 G; 8 T; 0 U; 0 Other;
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21-SEE-2000; 2000US-0234687P.
27-SEE-2000; 2000US-0236359P.
04-OCT-2000; 2000US-0236359P.
30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
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ABN08905 standard; DNA; 17 BP
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                                                                                                                                                                                                                                           17 ATGAGATGGACACAG 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                         RESULT 1460
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be expressing the proteins. The hGDMLP-1 proteins or polypeptides may be and/or amount specifically of hGDMLP proteins as specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser describion ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and in vaccines of for replacement therapy. The disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

Disclosure; SEQ ID NO 66; 214pp; English.

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protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as the rapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production with the expression of hGDMLP-1 in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in alectronic format directly from WIPO. It fip.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 8.7e+02; Matches 14; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 2 A; 8 C; 5 G; 2 T; 0 U; 0 Other;
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10-JAN-2001; 2001WO-US000664.
10-JAN-2001; 2001WO-US000665.
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30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000662.
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0-JAN-2001; 2001WO-US000668.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    165 ACTCCGAGGTGGCCG 179
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used an genome therspy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser describing in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and sequence encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 may be used for chromosome 22.

The present sequence represents an oligomer used in the screening of the horselfication, but was obtained in electronic format directly from WIPO capture, producincy production, but was obtained in electronic format directly from WIPO captured.
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or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                           Disclosure; SEQ ID NO 68; 214pp; English
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30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US000663.
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30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
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27-SEP-2000; 2000US-0236359P
04-OCT-2000; 2000GB-00024263
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30-JAN-2001; 2001WO-US000668
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 clan be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 mucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically of hGDMLP proteins, as specifically of hGDMLP proteins, as specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser described in onisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The adjaced sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders, hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. the protein but the present disorders but dispersion of the present invention. With the present sequence in the screening of the present sequence and the present disorders but and patent did not form part of the printed but the present sequence of the printed patent did not form part of the printed patent find patent find not form part of the printed patent find patent
                                                                                                                                                                                                                                                                                                                                                New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 8896; 214pp; English
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001US-0266860P.
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                                                                                                                                        (AEOM-) AEOMICA INC.
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. specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence 0; Gaps / Match 0.8%; Score 13.4; DB 1; Length 17; Local Similarity 93.3%; Pred. No. 8.7e+02; Les 14; Conservative 0; Mismatches 1; Indels Sequence 17 BP; 2 A; 7 C; 6 G; 2 T; 0 U; 0 Other; Query Match Matches

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ABO63456 standard; DNA; 17 BP RESULT 1463 ABQ63456/

20-AUG-2002 ABQ63456;

Human KTOMla portion (ABQ63232) probe # 169.

(first entry)

Human, KTOMIa; KTOMI, kidney tumour overexpressed membrane, cytostatic; gene therapy; cancer; kidney, liver, bone marrow, brain; heart; lung; kidney; colon; skeletal muscle; testis; uterus; placenta; probe; ss.

Homo sapiens

WO200224750-A2.

28-MAR-2002.

21-SEP-2001; 2001WO-US029656.

21-SEP-2000; 2000US-0234687P.

2001WO-US000665. 2001WO-US000666. 2001WO-US000667. 2001WO-US000669. 2001WO-US000669. 23-MAY-2001; 2001US-00864761. 28-AUG-2001; 2001US-0315676P. 2001WO-US000664 30-JAN-2001; 2 30-JAN-2001;

Shannon ME;

Chen W,

Hanzel DK, Rank DR,

Penn SG,

(AEOM-) AEOMICA INC.

Zhang J;

WPI; 2002-479509/51.

New human kidney tumor overexpressed membrane (KTOM1) protein and nucleic acids encoding the protein, useful for treating subjects having defects in KTOM1 which can manifest as cancer of the kidney, or as a disorder of e.g., liver or bone.

Example 2; Page 179; 418pp; English.

The invention relates to a novel isolated nucleic acid encoding human KTOM1 (kidney tumour overexpressed membrane) protein. The protein of the invention has cytostatic activity. The nucleotide may have a use in gene therapy. The KTOM1 nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOM1. Compositions comprising the nucleic acids, proteins or antibodies may be used to treat subjects having defects in KTOM1 which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, colon, skeletal muscle, testis, uterus and placenta function. The sequence represents a probe used in the invention to scan the nt 1-1001 portion of human KTOM1a (ABQ63232)

Sequence 17 BP; 5 A; 7 C; 3 G; 2 T; 0 U; 0 Other;

Gaps ö 0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; ative 0; Mismatches 1; Indels 14; Conservative Query Match Best Local Similarity Matches

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1397 AGCIGITGCAGTITG 1411 16 Agcrerrecaerere

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RESULT 1464 ABQ63457

ABQ63457 standard; DNA; 17 BP ABQ63457;

20-AUG-2002 (first entry)

Human KTOMla portion (ABQ63232) probe # 170.

Human; KTOMla; KTOMl; kidney tumour overexpressed membrane; cytostatic; gene therapy; cancer; kidney; liver; bone marrow; brain; heart; lung; kidney; colon; skeletal muscle; testis; uterus; placenta; probe; ss.

Homo sapiens.

WO200224750-A2.

21-SEP-2001; 2001WO-US029656.

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The present invention relates to human testis expressed Patched like
protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL
has two isoforms, with a few single base pair differences between the
two. One of the single base pair changes introduces a premature stop
codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL
codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL
codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL
codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL
codon in HTPL-S (S for short) stop in HTPL plays a role similar
consideration and second second second second second second second to the similar
consideration male germ cell development, and the HTPL gene was
mapped to human chromsome 10912.1. HTPL and its coding sequence are
useful for diagnosing a disorder caused by mutation in HTPL, and in
therapy and manufacture of a medicament for treatment or prevention of
therapy and manufacture of a medicament for treatment or prevention of
therapy and manufacture of a medicament for treatment or prevention of
therapy and manufacture of a medicament for treatment or prevention of
therapy and manufacture of a medicament for adrenal, adult and
foretal liver, bone marrow, brain, kidney, lung, placenta, prostate,
skeletal muscle or colon function. HTPL proteins and mucleic acids are
clinically useful diagnostic markers and potenial therapeutic agents for
male infertility and cancer. The present oligonucleotide was used in an
example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated human testis expressed Patched like protein (HTPL), ussful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
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                                                                         30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US0006664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001US-00864761.
09-OCT-2001; 2001US-00864761.
           28-JAN-2002; 2002EP-00001167
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           New human kidney tumor overexpressed membrane (KTOM1) protein and nucleic acids encoding the protein, useful for treating subjects having defects in KTOM1 which can manifest as cancer of the kidney, or as a disorder of e.g., liver or bone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel isolated nucleic acid encoding human KTCM1 (kidney tumour overexpressed membrane) protein. The protein of the invention has cytostatic activity. The nucleotide may have a use in gene therapy. The KTCM1 nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTCM1. Compositions comprising the nucleic acids, proteins or antibodies may be used to treat subjects having defects in KTCM1 which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, colon, skeletal muscle, testis, uterus and placenta function. The sequence represents a probe used in the invention to scan the nt 1-1001 portion of human KTCM1a (ABQ63232)
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-023555P.
04-OCT-2000; 2000US-023555P.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
23-WAY-2001; 2001WS-US000669.
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Gaps
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              0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                     ABV78819 standard; DNA; 17 BP
Query Match

Query Match

Best Local Similarity 93.5%,

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"shea 14; Conservative
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Gaps ; 0

Human, gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human teetis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss. Human HTPL scanning oligonucleotide SEQ ID 65.

Human, gene therapy, tumour suppressor; HTPL, chromosome 10p12.1; human testis expressed Patched like protein, testis, adrenal; liver; male germ cell development; bone marrow, brain, kidney, lung; placenta; prostate, skeletal muscle; colon, male infertility, cancer; ss.

EP1229046-A2 Homo sapiens

07-AUG-2002

Human HTPL scanning oligonucleotide SEQ ID 62.

(first entry)

03-JAN-2003

ABV78816;

ABV78816 standard; DNA; 17 BP

ò 원 03-JAN-2003 (first entry)

Homo sapiens

EP1229046-A2

Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;

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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV8762 and ABB9819 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structures strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10pl2.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in the useful for diagnosing a disorder caused by mutation in HTPL, and in the useful for diagnosing disorder caused by mutation of such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic agids are called infarility and cancer. The present oligomucleotide was used in an entered in an en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 0 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                               30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAX-2001; 2001US-00864761.
09-OCT-2001; 2001US-0327898P.
                                                  28-JAN-2002; 2002EP-00001167
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                                                                                                                                                                                                                                                                                                                                                   (AEOM-) AEOMICA INC.
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Best Local Similarity
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                                                                                                           30-JAN-2001;
07-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                      Zhan J;
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Matches
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The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, temporally, macular degeneration, the tumour angiogenesis, diabetic retinopathy, macular degeneration, necvascular glaucoma, myopic degeneration, arthritis, psoriasis, vertuce valaries, subjective solutions of the special portions of the relation of the patient having a condition associated with the level of ERG, by contacting a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies of the treatment of the conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treatment. (I) is useful for reducing ERG activity in a cell. by contecting the cell with (I). (I) is useful for cleaving RNA of the ERG gene, by contecting the cell with (I). (I) is useful for cleaving RNA of diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically trageting genes that share homology with ERG gene or ERG fusion genes. ABRZ7319 represent nucleic acids, including antisense and cenzymatic nucleic acid molecules which regulate expression of ERG, and cenzymatic nucleic acid molecules which regulate expression of ERG, and
                                                                                                                                                                                                                                                                                                                                                                                                                      Novel polynucleotide which down regulates expression of Bts-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
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0.8%; Score 13.4; DB 1; Length 17;
Best Local Similarity 80.0%; Pred. No. 8.7e+02;
Matches 12; Conservative 2; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 124; 149pp; English.
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                                                                                                                                                                                        16-MAY-2001; 2001WO-US015866.
                                                                                                                                                                                                                                  16-MAY-2000; 2000US-00572021.
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                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.
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                                                                                                    WO200188124-A2.
                                                             Homo sapiens.
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                                                                                                                                               22-NOV-2001.
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                      amberzyme.
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This invention describes a novel isolated nucleic acid that encodes one of three new isoforms of human pregnancy associated plasma protein E, harbap-E. The products of the invention have abortive and contraceptive activity and can be used for generation to in a vaccine. The nucleic acid, polypeptide encoded by it, or antibody to the polypeptide on be used in pharmaceutical compositions or vaccines for preventing or aborting pregnancy. PAPP-E is used in the antenatal diagnosis of dysgenetic pregnancies. The nucleic acids are used as probes to assess the level of PAPP-E isoform mRNA in chorionic villus samples, and the antibodies can be used to assess the expression levels of PAPP-E isoform proteins in chorionic villus samples, to diagnose dysgenetic pregnancies antenatally. This sequence represents an oligomer used in scanning the human PAPP-E genes described in the disclosure of the invention
PAPP-E; human; pregnancy associated plasma protein E; abortive; contraceptive; gene therapy; vaccine; pregnancy; antenatal; diagnosis; dysgenetic pregnancy; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPane; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid encoding an isoform of human pregnancy associated plasma protein E, for preventing or aborting pregnancy
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                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 147; 353pp; English
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(SHAN/) SHANNON M E.
                                                                                                                                                                                                                                                                                                                                      WPI; 2002-697817/75.
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                                                                                                     US2002102252-A1.
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                                                                     Homo sapiens.
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSH2) in polypeptide (1), comprising a sequence of 730 amino acids (SI, ABB8399), a sequence having 65% sequence identity to (SI), acids (SI) having 95% deviations, especially conservative substitutions or a graph of the sequences comprising at least 8 contiguous amino acids. Human POSH1 1 is a proto-oncogene/Oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (II) is useful considered by altered expression of human POSH1 including diagnosing and creating cancer, they useful in the development of vaccines and treating treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonuclectide useful in examples of free invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to betwent by the Buropean Patent Office
                                                                                                                                                                                                                                                                                                                   Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, FOSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; SEQ ID NO 977; 60pp + Sequence Listing; English.
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                              30-JNN-2001; 2001WO-US000665.
30-JNN-2001; 2001WO-US000666.
30-JNN-2001; 2001WO-US000667.
30-JNN-2001; 2001WO-US000669.
30-JNN-2001; 2001WO-US000669.
23-MAX-2001; 2001WG-US000670.
10-OCT-2001; 2001US-0328205P.
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Matches 14; Conservative
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                                                                                                                                                                                                        (AEOM-) AEOMICA INC.
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (SL, ABB83999), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring dispease and treating cancer, they useful in the development of vaccines and (II) is cuseful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to betwent by the Buropean Patent Office
                                                                                                                                                                                                                                                                                                                                                         Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
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                                                                                                                    30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WS-US000670.
23-MAY-2001; 2001US-00864761.
10-CCT-2001; 2001US-0328205P.
                                                                  30-JAN-2001; 2001WO-US000065.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
                   28-JAN-2002; 2002EP-00001165
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Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.

30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000666. 30-JAN-2001; 2001WO-US000667. 30-JAN-2001; 2001WO-US000669. 30-JAN-2001; 2001WO-US000669. 23-YAN-2001; 2001WG-US000670. 23-YAX-2001; 2001US-0328205P.

(AEOM-) AEOMICA INC.

WPI; 2002-684061/74.

Shannon M;

28-JAN-2002; 2002EP-00001165

EP1239051-A2 11-SEP-2002. Example 2; SEQ ID NO 1806; 60pp + Sequence Listing; English.

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ABV90265;
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Rho GTPase; signal transduction; gene expression; cancer; vaccine;
gene therapy; transgenic; ss.
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBE3999), a sequence having 65* sequence identity to (SI), (G1) having 95* deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating encoding (I) are useful for diagnosing monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence difformation supplied to printed specification, but is based on sequence information supplied to browny and the present of the content of the present of the content of the present of the content of the
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Rho GTPase; signal transduction, gene expression; cancer; vaccine;
gene therapy; transgenic; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 8.7e+02; ative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human POSHL1 scanning oligonucleotide SEQ ID NO 978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 7 A; 2 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1451 ATCCATTCTTCCTCA 1465
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Best Local Similarity 93.3%
Matches 14; Conservative
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AC ABV9
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DT 23-I
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KW Hume
KW Rho
KW Rho
KW Gent
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ABV90266
ID ABV90266 standard; DNA; 17 BP.
XX
AC ABV90266;
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DT 23-DEC-2002 (first entry)
XX
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DE Human POSHL1 scanning oligonucle
XX
                                                                                       30-JAN-2001; 2001WO-US000670.
23-MAY-2001; 2001US-00864761.
10-OCT-2001; 2001US-0328205P.
                                     28-JAN-2002; 2002EP-00001165
                                                                                                                                    WPI; 2002-684061/74.
                                                                                                             (AEOM-) AEOMICA INC
                                                      30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
      Homo sapiens.
                EP1239051-A2
                                                 30-JAN-2001;
                           11-SEP-2002
                                                                                                                         Shannon M;
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Matches
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (61, ABB83999), a sequence having 65% sequence of 631), (61) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPaces as well as downstream components of the signal transduction pathway. (I) is useful cfor identifying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and creating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating cance useful in measuring and for surveying gene expression and creating present sequence is that of a scanning oligonuclectide useful in examples of the invention. Note: The present sequence did not form part of the content the provent of the content of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, FOSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
   Human, POSHL 1; SH3 domain, POSH-like signalling protein 1; oncogene, Rho GTPase; signal transduction, gene expression, cancer; vaccine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JAN-2001; 2001WO-US006667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
23-MAX-2001; 2001WG-0800670.
                                                               gene therapy; transgenic; ss.
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ABV91091 standard; DNA; 17 BP.
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30-JAN-2001;
                                                                                                                               Homo sapiens,
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ABV91091/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
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2001WO-US000668.
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